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Result
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Maximum DB
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                                                                                                                                                                                                                       Score
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length: 2000000000
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1: geneseqp1980s:*
2: geneseqp1990s:*
3: geneseqp2000s:*
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AAM24521
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AAW84274
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Sequence 158 AA;

Claim 1; Page

53; 64pp; English.

A human colon specific protein (AAW12691) is a potential diagnostic marker for colon cancer. It is believed that the presence of active transcription of the colon specific gene in non-colon cells of a host is indicative of colon cancer metastases. The amino acid sequence of the colon specific protein was deduced from a cDNA clone (AATS1784) isolated from a human colon cancer cDNA library. Recombinant colon specific protein can be produced in transformed host (e.g. bacterial, insect) cells and used to develop prods. for the diagnosis and treatment of colon disorders, partic. colon cancer metastasis. Antibodies raised against the protein can be used to target cancer cells and as part of a colon cancer

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New isolated diagnosis an	-1997-04 DB; AATS	et DR,		06-JUN-1995	06-JUN-1995	12-DEC-1996	WO9639541-A1	sapiens	spec	Human colon	31-MAY-1997	AAW12691;	AAW12691 81			254.5	307.5	760	856	867	878 878	878	878	878	878 878	878	878	878 878	878	878
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prods. for use i colon cancer.				٠					diagnosis; therapy;							1513	Abp59097 Human			Abb99310 Amino		Aag75620 Human			Adi58211 Human		-	Adn39882 Cancer	72	Adf85474 Human
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                                                                                                                                                                      The present sequence represents a human Reg I-gamma protein, which comprises a C-type lectin. The sequence was identified in Incyte clone 1310334. Reg I-gamma protein is involved in regulation of cell growth and development. Since the overexpression of reg proteins is associated with tumourogenesis and neurodegeneration, inhibition of human Reg I-gamma expression can be used for treating or preventing neoplasia or metastasis and neurodegenerative changes associated with Alzheimer's disease and characters of the catalogs associated with Alzheimer's disease and catalogs as a sequence was a sequence which are the catalogs as a sequence was a sequence w
                                                                     other disorders of the central nervous system, e.g. Down's syndrome. Reg I-gamma can also be used in therapeutics to induce regeneration of pancreatic beta-cells in the treatment of diabetes. The products can also be used for detection for, e.g. expression of REG I-gamma, diagnosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; Reg I-gamma protein; C-type lectin; Incyte clone 1310334; regulation; cell growth; development; tumourogenesis; neurodegeneration; inhibition; treatment; prevention; neoplasia; metastasis; neurodegenerative change; Alzheimer's disease; Down's syndrome; regeneration; pancreatic beta-cells; diabetes.
                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Fig 1; 72pp; English
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Query Match

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                                                                                                                                                                                                                                                                                     This is the amino acid sequence of a novel human protein comprising a secretory signal isolated from stomach cancer cells. Its proteins can used as nutritional sources or supplements. The proteins may also have cytokine functions, immune modulating functions, haematopoiesis regulating activity, activin/inhibin regulating activity, chemotactic/chemokinetic activity, haemostatic and thrombolytic activity receptor/ligand activity, anti-inflammatory activity, tumour inhibitio
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                                              MASRSMRLLLLLSCLAKTGVLGDIIMRPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQS
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  MASRSMRLLLLLSCLAKTGVLGD1 IMRPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQS
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                                                                                                          Conservative
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                                                                                                                             100.0%;
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                                                                                                                               Score 878; DB 2; Pred. No. 6.8e-83;
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                                                                                                          Mismatches
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                                                                                                                                                        Length 158;
                                                                                                       Indels
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                                                                                                                                                 Query Match
Best Local S
Matches 158
                                                                                                                                                                                                                       The present sequence is encoded by a human colon specific gene. The nucleic acid sequence can be used to develop products for the diagnosis of a disorder of the colon, e.g. colon cancer or metastases. The product can also be used to screen for agonists or antagonists for the polypeptides. The antagonists may be used to treat colon cancer, since they interact with the function of colon specific polypeptides to inhib functions which are necessary for the viability of colon cancer cells. The products can also be used for the production of antibodies and for the identification of receptors for the polypeptides
                                                                                                                                                                                                  Sequence 158 AA;
                                                                                                                                                                                                                                                                                                                                                   Claim 1; Fig 1A-C; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                metastases.
                                                                                                                                                                                                                                                                                                                                                                                         Isolated human colon specific gene - diagnosis and treatment of disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                 N-PSDB; AAX03195.
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               KSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKYRP 158
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KSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKYRP 158
                                                YGNGAHLASILSLKEASTIAEYISGYQRSQPJWIGLHDPQKRQQWQWIDGAMYLYRSWSG
                                                                YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRSWSG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   a human colon specific gene.
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Pred. No. 6.8e-83;
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RESULT 6
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XX 127-J
XX 29-S
XX 19-S
XX WPI;
DR N-BS
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Best Local S
Matches 158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to a purified human protein, which is primarily expressed in tissue derived from the colon. The protein is 152 amino acids in length and exhibits cytostatic activity. The present sequence represents the amino acid sequence of the colon specific protein. The protein can be used in the diagnosis and treatment of colon cancer, and it is thought that abnormally high levels of the gene expression in non colon cells is an indication of colon cancer metastasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Fig 1; 20pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel human colon specific polypeptides and polynucleotides for diagnosis and treatment of colon cancer, for screening compounds which interact with polypeptide, for synthesis of DNA and manufacture of DNA vectors.
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                                  KSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKYRP
                                                                                                                          YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRSWSG
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KSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKYRP
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Pred. No. 6.8e-83;
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RESULT 7

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10-JAN-2000;
15-FEB-2000;
                                                                                                          29-DEC-2000; 2000WO-US035596.
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                                                                                                                                                                                                                WO200149716-A2
                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                             C880P similar amino acid sequence (GENESEQ W37866).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1;
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                                                                                                                                                                                                                                                                                                                therapy;
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                                                                                                                                                                                                                                                                                                                                     immunotherapy; diagnosis; colon cancer; colon tumour; immunogenic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 MASRSMRLLLLLLSCLAKTGVLGDIIMRPSCAPGMFYHKSNCYGYFRKLRNWSDAELECQS
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  99US-00476296.
2000US-00480321.
2000US-00504629.
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                                                                                                                                                                                                                                                                                                                     vaccine;
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                                                                                                                                                                                                                                                                                                                     colonic
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Pred. No. 6.8e-83;
                                                                                                                                                                                                                                                                                                                     cancer.
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RESULT 8
AAM24520
ID AAM2
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AC AAM2
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DT 12-C
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DT 12-C
XX
DT U2-C

(first entry) amino

AAM24520 standard; protein; 158 AA

gene therapy;

C880P similar 12-OCT-2001

acid

sequence

(GENÉSEQ W37929).

Human; immunotherapy; diagnosis; Colon cancer; colon tumour; immunogenic; gene therapy; vaccine; colonic Cancer.

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Best Local (
                                                                                                                                                                                                                                                                                                                                                               TCAPS by expressing inactive proteins or to supplement the patients own production of them. Additionally, (II) may be used to produce the TCAP proteins, by inserting the nucleic acids into a host cell culturing the cell to express the protein. (II) and its complementary sequences may also be used as DNA probes in diagnostic polymerase chain reaction (PCR) and hybridisation assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. (I) may also be used as antigens in the production of antibodies against TCAPs and in assays to identify modulators of TCAP expression and activity. Anti-(I) antibodies and antagonists may also be used as diagnostic agents for detecting the presence of TCAPs in samples (e.g. by enzyme linked immunosorbant assay (BLISA)). AAI28460 to AAI28512 and AAM24494 to AAM24523 represent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-MAR-2000;
19-MAY-2000;
29-JUN-2000;
28-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            associated with inappropriate colon tumour associated protein (TCAP) expression, such as colonic cancer. For example, (I) and (II) may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               χυ,
                                                                                                                                                                                                                                                                                             Sequence 158 AA;
                                                                                                                                                                                                                                                                                                                                                     nucleotide and amino
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Colon tumor associated proteins and nucleic acids useful prevention, diagnosis and treatment of colonic cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-441847/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes colon tumour associated proteins (I) and the polynucleotides (II) that encode them. (I) have cytostatic activity. (I) and (II) can be used in gene therapy and vaccine production. (I) and (II) may be used in the prevention, diagnosis and treatment of diseases
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121
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E, Wang T,
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                                                                                                                                                                                                                                                                                                                                  invention
                                                                                                 YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRSWSG 120
                                KSMGGNKHCAEMSSNNNFLTWSSNECNKROHFLCKYRP 158
KSMGGNKHCAEMSSNNNFLTWSSNECNKROHFLCKYRP 158
                                                                      YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRSWSG
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                                                                                                                                                                                                                    Conservative
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; 2000US-00609448.
; 2000US-00649811.
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Jiang Y;
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                                                                                                                                                                                                                                        Score 878; DB 4;
Pred. No. 6.8e-83;
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Best Local
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15-FEB-2000;
06-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Colon tumor associated proteins and nucleic acids useful for the prevention, diagnosis and treatment of colonic cancer.
                                                                                                                                                                                                                                                                                               Sequence 158 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; Page 468; 472pp; English.
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29-JUN-2000;
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B
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121
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E, Wang T,
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                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                    invention
                     KSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKYRP 158
                                                                                                     YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRSWSG
KSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKYRP 158
                                                                       YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRSWSG
                                                                                                                                               MASRSMRLLLLLSCLAKTGVLGDIIMRPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQS
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2000US-00504629.
2000US-00519444.
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2000US-00609448.
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Jiang Y;
                                                                                                                                                                                                                     100.0%; Score 878; DB 4;
100.0%; Pred. No. 6.8e-83;
htive 0; Mismatches 0;
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                                                                                                                                                                                                                                                        Length 158;
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the polynucleotides (II) that encode them. (I) have cytostatic activity. (I) and (II) can be used in gene therapy and vaccine production. (I) and (II) may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate colon tumour associated protein (TCAR) and (II) may be used in the prevention, diagnosis and treatment of diseases (II) at the colonic cancer. For example, (II) and (II) may be used to treat disorders associated with decreased expression by rectifying (II) the colonic cancer of the catherity of the colonic cancer of the catherity of the cathering the nucleic acids in the patients own (II) may be used to produce the TCAR (II) and its complementary sequences may (II) to express the protein. (II) and its complementary sequences may (II) be used as DNA probes in diagnostic polymerase chain reaction (PCR) and hybridisation assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of capression and activity. Anti-(I) antibodies and antagonists may also be used as antigens in the production (C antibodies may also be used as antigens in the production (C capression and activity. Anti-(I) antibodies and antagonists may also be used as antibodies and antagonists may also be used as antibodies and antagonists may also be used as diagnostic agents of the anti-(I) antibodies and activity. The anti-(I) antibodies and antibodie
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06-MAR-2000;
19-MAY-2000;
                                                                                          Sequence 158 AA;
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                                                                                                                                      present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; Page 469; 472pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Colon tumor associated proteins and nucleic acids useful prevention, diagnosis and treatment of colonic cancer.
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3E, Wang T,
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Jiang Y;
Score 878; DB 4;
Pred. No. 6.8e-83;
D: Mismatches 0;
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Query Match
Best Local Similarity
Matches 158; Conserv

Conservative

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Length 158; Indels

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AAM24517
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                  The present invention describes colon tumour associated proteins (I) and (II) can be used in gene therapy and vaccine production. (I) and (II) can be used in gene therapy and vaccine production. (I) and (II) may be used in the prevention, diagnosis and treatment of diseases (II) may be used in the prevention, diagnosis and treatment of diseases casociated with inappropriate colon tumour associated protein (TCAP) expression, such as colonic cancer. For example, (I) and (II) may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of TCAPS by expressing inactive proteins or to supplement the patients own production of them. Additionally, (II) may be used to produce the TCAP proteins, by inserting the nucleic acids into a host cell culturing the cell to express the protein. (II) and its complementary sequences may call to express a DNA probes in diagnostic polymerase chain reaction (PCR) and hybridisation assays to detect and quantitate the presence of similar nucleic acids in samples, and therefore which patients may be in need of restorative therapy. (I) may also be used as antiponists may also be expression and activity. Anti-(I) antibodies and antagonists may also be used to down regulate TCAP expression and activity. The anti-(I) antibodies may also be used to down regulate TCAP expression and activity. The anti-(I) antibodies may also be used to down regulate TCAP expression and activity. The anti-(I) antibodies may also be used to down regulate TCAP expression and activity. The anti-(I) antibodies may also be used to down regulate TCAP expression and activity.
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29-JUN-2000;
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15-FEB-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Lodes MJ
E, Wang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  immunotherapy; diagnosis; colon cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 463; 472pp; English.
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2000US-00480321.
2000US-00504629.
2000US-00519444.
2000US-00575251.
2000US-00609448.
2000US-006094881.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Secrist H,
Jiang Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            colonic
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as diagnostic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acids useful of colonic cancer.
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   agents
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     colon tumour; immunogenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            158
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   for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Stolk JA;
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                                                                                                                                                                                                                                                         06-MAR-2000;
19-MAY-2000;
29-JUN-2000;
                                                                                                                                                                                      Xu J,
King
                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; immunot gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        presence of TCAPs in samples (e.g. by enzyme linked immunosorbant assay (ELISA)). AAI28460 to AAI29512 and AAM24494 to AAM24523 represent nucleotide and amino acid sequences given in the exemplification of the
                                         The present invention describes colon tumour associated proteins (I) and the polynucleotides (II) that encode them. (I) have cytostatic activity. (I) and (II) can be used in gene therapy and vaccine production. (I) and (II) may be used in the prevention, diagnosis and treatment of diseases
                                                                                                       Claim 2;
                                                                                                                            prevention,
                                                                                                                                                              WPI; 2001-441847/47
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                                                                                                                                                                                                                                                28-AUG-2000;
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15-FEB-2000;
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                                                                                                                                                                                                                         (CORI-) CORIXA CORP.
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                                                                                                                                       tumor
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                                                                                                                                                                                    Lodes MJ,
3E, Wang T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                     immunotherapy; diagnosis; colon cancer; colon tumour; immunogenic;
herapy; vaccine; colonic cancer.
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                                                                                                       Page
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                                                                                                                            associated proteins and diagnosis and treatment
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2000US-005194629.

2000US-00519444.

2000US-00575251.

2000US-00609448.

2000US-00649811.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                     467; 472pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid sequence (GENESEQ W12691).
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                                                                                                                                                                                    Secrist H,
Jiang Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                       English.
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Pred. No. 6
                                                                                                                                                                                                  Benson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                              nucleic acids useful of colonic cancer.
                                                                                                                                                                                                  В,
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                                                                                                                                                                                                  Meagher
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associated with inappropriate colon tumour associated protein (TCAP) expression, such as colonic cancer. For example, (I) and (II) may be used to treat disorders associated with decreased expression by rectifying mutations or deletions in a patient's genome that affect the activity of

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RESULT 12
AAE29829
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Best Local Similarity
Detecting the presence of a tumor comprises detecting the concentration of a Reg Like Protein or the presence or quantity of a nucleic acid
                                                                                 WPI; 2002-684095/74.
N-PSDB; AAD47239.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Key
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                                                                                                                                                                      Heiskala M;
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                                                                                                                                                                                                                           CLINICAL DIAGNOSTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Human mature REG-like protein"
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Pred. No. 6.8e-83;
): Mismatches 0;
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method for detecting REG-like protein (RELP) and its nucleic acid sequence. The method is useful for detecting the presence of a tumour. Kits comprising an antibody specific for RELP and reagents for detecting the antibody, or a nucleic acid complementary to portion of a nucleic acid encoding RELP, are useful for identifying the presence of cancer, characterise the cancer, or monitor the course of treatment of cancer. The present sequence is human RELP protein used to illustrate the method of the invention. Human RELP gene is located at chromosome 1
                                                                                                                                                                                                                                                                                                                                                                                                                       Human; colon cancer; cancer; tissue profiling; forensic; genetic analysis; diagnostic; antisense therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 158 AA;
ABQ56306 to ABQ60787 represent isolated nucleic acids (I) differentially expressed in cancer tissues. ABB78993 to ABB79004 represent proteins encoded by the ABQ60776 to ABQ60787 nucleic acid sequences. (I) can be
                                                                                    New isolated nucleic acid that is differentially expressed in cancer tissues useful for determining the presence of colon cancer in a cell or tissue type, and in antisense therapy.
                                                                                                                                                                                          Burgess C, Astle JH,
Thiaglingam A, Lewis
                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human Reg IV protein sequence SEQ ID NO:4471.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-AUG-2002
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                                                           Claim 5;
                                                                                                                                                   N-PSDB;
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DB; ABQ60776.
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                                                           Fig 3; 796pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKYRP 158
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                                                                                                                                                                                               Lewis
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RESULT 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           used in antisense therapy. An antibody immunoreactive with a polypeptide encoded by (I) is useful for detecting cancer in a patient sample, and for detecting the presence or absence of a polynucleotide encoded by a nucleic acid which hybridises to (I) in a cell. A probe/primer derived from (I) can be used for determining the presence of a nucleic acid which hybridises to (I), and for determining the phenotype of cells in a sample of cells from a patient. (I) is useful for determining the presence of colon cancer in a cell or tissue type, for determining the presence or state of other type of cancer, in antisense therapy, to generate macroarrays on a solid surface, to identify a chromosome on which the corresponding gene resides, and in tissue profiling, forensics, genetic analysis, mapping and diagnostic applications. (I) can be used to raise antibodies, and to screen for peptide analogues and antagonists
New genes that are up-regulated or down-regulated in cancers, useful as markers for diagnosing e.g. cancer, ischemia or heart diseases, or as therapeutic targets for screening drugs for treating these diseases.
                                                                                                                                                                                Zlotnik
                                                                                                                                                                                                    Afar D,
                                                                                                                                                                                                                                                                                                   08-FEB-2002;
12-APR-2002;
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13-NOV-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human cancer related protein SEQ ID NO:208.
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                                                                                                      2003-354600/33.
DB; ACC72672.
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                                                                                                                                                                                                                                                                                              ; 2001US-0323469P.
; 2001US-0353887P.
; 2001US-0350666P.
; 2002US-0355145P.
; 2002US-0355257P.
; 2002US-0372246P.
                                                                                                                                                                                                                                                 BIOTECHNOLOGY INC
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                                                                                                                                                                                                    Gish KC,
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Pred. No. 6.8e-83;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes an isolated nucleic acid molecule, which CC comprises the sequence of any of the genes that are up-regulated or down-cc regulated in specific cancers (e.g. about 1031 genes up-regulated in CC acute lymphocytic leukemia). ACC72641 to ACC72860 represent cancer cc and the lymphocytic leukemia). ACC72641 to ACC72860 represent cancer cc absence of a pathological cell in a patient; (2) an expression vector cc comprising a nucleic acid molecule described: (1) determining the presence or cc comprising the vector; (4) an isolated polypeptide, which is encoded by cc comprising the vector; (5) an antibody that specifically binds the polypeptide cc of (4); (6) specifically targeting a compound to a pathological cell in a cc cross screening assay. The nucleic acid is useful as diagnostic markers or therapeutic targets. In particular, the nucleic acid is useful for cd diagnosing a pathology, e.g. cancer (e.g. cancer of the bone marrow, cc bladder, brain, breast, crevix, colon/rectum, kidney, lung, ovary, cc pathologies and endometriosis. The nucleic acid is also useful in constitution and uterus), wounds, ischaemia, heart diseases, constablories.
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Matches
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                                                           16-MAR-2001; 2001US-0276305P
                                                                                                                      14-MAR-2002; 2002WO-US007945
                                                                                                                                                                                     26-SEP-2002.
                                                                                                                                                                                                                                                                                                                                     Protein
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                                                                                                                                                                                                                                                                                                     /label= signal
27. .158
/label= RELP
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Pred. No. 6.8e-83;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID
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Matches 158
                                                                                                                                                                                                                                                                                                    The present sequence represents a new isolated REG-like protein (RELP) human immunoglobulin (Ig) derived protein. RELP comprises: (a) a human variable and constant region; or (b) an isolated human Ig derived protein or specified portion or variant encoded by a nucleic acid. RELP has cytostatic activity and can be used as an Ig agonist and in protein therapy. The RELP human Ig derived protein or a specified portion or variant can be used for preventing or treating a RELP protein mediated condition, malignant condition or disease condition, e.g. cancer. The nucleic acids can be used in producing RELP Ig derived protein. The human RELP protein of the present invention is located to chromosome 1p12-13.1
                                                                                                                                                                                                                                                                         Sequence 158 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Fig 2; 101pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated REG-like protein (RELP) human immunoglobulin derived protein or specified portion or variant, useful for preventing or treating a RELP protein mediated condition or malignant condition, e.g. cancer.
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N-PSDB; ABZ21635.
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nes 158; Conserv
                 121 KSMGGNKHCAEMSSNNNPLTWSSNECNKRQHFLCKYRP 158
121
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                                                                                                                              KSMGGNKHCAEMSSNNNFLTWSSNECNKROHFLCKYRP 158
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  score greater than or equal to the score of the result being and is derived by analysis of the total score distribution.
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2: uniprot_trembl:*
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Q9D8G5
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Q6qx33
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RESULT 1
Q9BYZ8
         Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

Al Klausner R.D., Collins F.S., Wagner L., Schemmen C.M., Schuler G.D.,

Al Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Botak S.S., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carminci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gunaratne P.H.,

Carmino J., Hulyk S.W.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Van Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Krzywinski M.I., Skalska U., Smailus D.E., Schmerch A., Schein J.E.,

Jones S.J., Marra M.A.,

Jones S.J., Marra M.A.,

Jones S.J., Marra M.A.,
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Q9BYZ8;
01-JUN-2001
01-JUN-2001
25-OCT-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.

MEDILINE=21210973; PubMed=11311942; DOI=10.1016/S0167-4781(00)00284-0;
Hartupee J.C., Zhang H., Bonaldo M.F., Soares M.B., Dieckgraefe B.K.;
"Isolation and characterization of a cDNA encoding a novel member of
the human regenerating protein family: Reg IV(1).";
Biochim. Biophys. Acta 1518:287-293(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2001 (TrEMBLrel. 28, Last annotation update)
25-OCT-2004 (TrEMBLrel. 28, Last annotation update)
Regenerating gene type IV precursor (REG-like protein) (Regenerating islet-defived family, member 4) (Gastrointestinal secretory protein
                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
Kamarainen M., Heiskala
Submitted (JUN-2002) to
                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Colon cancer;
Violette S., Festor E.,
Lacasa M., Lesuffleur T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Name REG4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GISP)
                                                                                                                                                                                                                                                                       MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                 Submitted (FEB-2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
Generation and initial analysis of more than 15,000 full-length humar
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                                                                                                                                                                                                                                                                                                                                      Heiskala M., Andersson L.C.;
EMBL/GenBank/DDBJ databases.
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Q6X5S3
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11 homo sapien
12 echis pyram
3 echis ocell
9 echis carin
1 bitis ariet
0 echis multi
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RIKEN FANTOM Consortium;

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RESULTI OPERACE OF ACCORD 
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Best Local S
Matches 158
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EMBL; AF345934, AAK59869.1;
EMBL; AY126670; AAM95598.1;
EMBL; BC017089; AAH17089.1;
EMBL; AF254415; AAK48435.1;
HSSP; P22030; IIJK.
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Q9D8G5;
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GO; GO:0005529; F:sugar binding; IE
InterPro; IPR001304; Lectin C.
InterPro; IPR003990; Pancreatis_ac.
                                        SEQUENCE FROM N.,
STRAIN=C57BL/6J;
                                                                                                 Carninci P., Hayashizaki Y.;
"High-efficiency full-length c
Meth. Enzymol. 303:19-44(1999)
                                                                                                                                                                                  SEQUENCE FROM N.A. STRAIN=C57BL/6J; TISSUE=Small intestine;
                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                     Mus musculus adult male enriched library, clone
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25-OCT-2004 (TrEMBLrel.
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PRINTS; PR01504; PNCREATITSAP.
SMART; SM00034; CLECT; 1.
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c. Natl. Acad. Sci. U.S.A.
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llarity 100.0%;
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.A.
/; TISSUE=Small intestine;
50; PubMed=11217851; DOI=1
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BLrel. 17, Last sequence update)
BLrel. 28, Last annotation update)
t male small intestine cDNA, RIKEN full-length
clone:2010002L15, product:REGENERATING GENE TY
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Rodentia;
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Pred. No. 2.5e-78;
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Sciurognathi;
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                                                                                                                    cloning.";
                     DOI=10.1038/35055500;
                                                                                                                                                               DOI=10.1016/S0076-6879(99)03004-9;
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chi; Muridae;
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WEDLINE-22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

& Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

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& Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

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& Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Mullahy S.J.,

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& Raha S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

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& Villalon D.K., Muzny D.M., Sodergen E.J., Lu X., Gibbs R.A.,

& William M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

& Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

& Balkesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

& Balkesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
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Arakawa T., Bono H., Carninci P., Fukuda S., Fukunishi Y., Furuno M.
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Sano H., Sasaki D., Shibata K., Shibata Y., Shinagawa A., Shiraki T.,
Sogabe Y., Suzuki H., Tagami M., Takahashi F., Tanaka T.,
Tejima Y., Toya T., Yamamura T., Yasunishi A., Yoshida K., Yoshino Muramatsu M., Hayashizaki Y.,
Muramatsu M., Hayashizaki Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN-C57BL/6J; TISSUB-Small intestine;

MEDLINB-20530913; PubMed=11076861; DOI=10.1101/gr.152600;

Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P.,

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Konno H., Akiyama J., Nishi K., Kitsunai T., Tashiro H., Itoh M.,

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Genome Res. 10:1757-1771(2000).
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SEQUENCE FROM N.A.

STRAIN=C57BL/6J; TISSUB=Small intestine;

STRAIN=C57BL/6J; TISSUB=Small intestine;

MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100;

MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100;
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                                                                                          Fahey J., Helton E., Ketteman M., Madan A., Rodrigue Whiting M., Madan A., Young A.C., Shevchenko Y., Gue Whiting M., Madan A., Young A.C., Shevchenko Y., Dickson N. Blakesley R.W., Touchman J.W., Green E.D., Dickson N. Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Krzywinski M.I., Skalska U., Smailus D.E., Schnerch
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Nature 409:685-690(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=FVB/N;
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                                                                        Jones S.J., Marra M.A.
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57BL/6J; TISSUE=Small intestine;
Akimur
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                    on and initial analysis cDNA sequences.";
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on functional annotation
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                                                                                                                            R.M.,
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                                                                                                                         Butterfield
                                                   full-length human
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Prange C.,
Mullahy S.J.,
Inaratne P.H.,
J., Hulyk S.W.,
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RESULT 3

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Best Local (
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01-JUN-2001
01-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-VAR-2001 (TrEMBLrel. 17, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Mus musculus adult male small intestine cDNA, RIESN full-length
enriched library, clone:2010204K21 product:REGENERATING GENE TYP
full insert sequence.
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SEQUENCE FROM N.A.
STRAIN=FVB/N; TISSUE=Colon;
There R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (DEC-2001) to the EMBL/GenBank/DDBJ databases. EMBL; AK008049; BAB25429.1; -. EMBL; BC019465; AAH19465.1; -. HSSP; Q06141; 1UV0.
                                           STRAIN=C57BL/6J; TISSUE=Small intestine; The PANTOM Consortium, the RIKEN Genome Exploration Research Gr "Analysis of the mouse transcriptome bas 60,770 full-length CDNAs.";
                                                                                                                                                                                                                                                                                                                                                                            STRAIN=C57BL/6J; TISSUE=Small intestine;
MEDLINE=99279253; PubMed=10349636; DOI=10.1016/S0076-6879(99)03004-9;
Carninci P., Hayashizaki Y.;
"High-efficiency full-length cDNA cloning.";
Meth. Enzymol. 303:19-44(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
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01-JUN-2001 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
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SMART; SM00034; CLECT; 1.
PROSITE; PS50041; C. TYPE LECTIN 2; 1.
SEQUENCE 157 AA; 18398 MW; F3981722BBD83968 CRC64;
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                                                                                                                                                                   SEQUENCE FROM
                                                                                                                                                                                                                  Nature 409:685-690(2001).
                                                                                                                                                                                                                                                              RIKEN FANTOM Consortium;
                                                                                                                                                                                                                                                                                    MEDLINE=21085660; PubMed=11217851;
                                                                                                                                                                                                                                                                                                            STRAIN=C57BL/6J;
                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (Mouse)
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                          420:563-573 (2002).
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                                                                                                                                                                                                                                     annotation of a full-length
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Rodentia;
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criptome based o
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Sciurognathi; Muridae;
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                                                                      on functional
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                                                                      Team;
annotation
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RESULT 4
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RC Adachi J., Aizawa K., Akahira S., Akimura T., Arai A., Aono H.,

RA Adachi J., Aizawa K., Akahira S., Akimura T., Arai A., Aono H.,

RA Adachi J., Aizawa K., Akahira S., Akimura T., Arai A., Aono H.,

RA Arakawa T., Bono H., Carninci P., Fukuda S., Fukunishi Y., Furuno M.,

RA Hanagaki T., Hara A., Hayatsu N., Hiramoto K., Hiraoka T., Kato H.,

RA Imotani K., Ishii Y., Itoh M., Izawa M., Kasukawa T., Kato H.,

RA Kawai J., Kojima Y., Konno H., Kouda M., Koya S., Kurihara C.,

RA Kawai J., Miyazaki A., Nahi K., Nomura K., Numazaki R., Ohno M.,

RA Mateuyama T., Miyazaki A., Nahi K., Nomura K., Numazaki R., Ohno M.,

RA Adachi J., Nidara K., Shibata Y., Shinagawa A., Shiraki T.,

RA Sano H., Sasaki D., Shibata K., Shibata Y., Shinagawa A., Shiraki T.,

RA Sogabe Y., Suzuki H., Tagami M., Tagawa A., Taskahashi F., Tanaka T.,

RA Tejima Y., Toya T., Yamamura T., Yasunishi A., Yoshida K., Yoshino M.,

RMIL, AKOO8438, BAB25669.1; -.

RMIL, AKOO8438, BAB25669.1; -.
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Matches 103
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MEDLINE=20530913, PubMed=11076861, DÖI=10.1101/gr.152600;

MEDLINE=20530913, PubMed=11076861, DÖI=10.1101/gr.152600;

Shibata K., Itoh M., Aizawa K., Nagaoka S., Sasaki N., Carninci P.,

Konno H., Akiyama J., Nishi K., Kitsunai T., Tashiro H., Itoh M.,

Yamamoto R., Matsumoto H., Sakaguchi S., Kegami T., Kashiwagi K.,

Yamamoto R., Matsumoto H., Sakaguchi S., Negami T., Kashiwagi K.,

Pujiwake S., Inoue K., Togawa Y., Izawa M., Ohara E., Watchiki M.,

Yoneda Y., Ishikawa T., Ozawa K., Tanaka T., Matsuura S., Kawai J.,

Yokazaki Y., Muramatsu M., Inoue Y., Kira A., Hayashizaki Y.;

"RIKEN integrated sequence analysis (RISA) system-384-format

sequencing pipeline with 384 multicapillary sequencer.";

Genome Res. 10:1757-1771(2000).
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PRINTS; PR01504; PNCREATITSAP.
SMART; SM00034; CLECT; 1.
PROSITE; PS50041; C_TYPE_LECTIN_2; 1.
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STRAIN-C57BL/6J; TISSUB=Small intestine;

MEDLINE=20499374; PubMed=11042159; DOI=10.1101/gr.145100;

Carninci P., Shibata Y., Hayatsu N., Sugahara Y., Shibata K.,

Konno H., Okazaki Y., Muramatsu M., Hayashizaki Y.;

"Normalization and subtraction of cap-trapper-selected cDNAs t
prepare full-length cDNA libraries for rapid discovery of new
fenome Res. 10:1617-1630(2000).
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157 AA; 18474 MW; FD96F36CFB989368 CRC64;
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Pred. No. 1.9e-50;
5; Mismatches 28;
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Q68AX7; Q68AX7; 25-OCT-2004 25-OCT-2004 25-OCT-2004

(TrEMBLrel. (TrEMBLrel. (TrEMBLrel.

28,

Created)
Last sequence update)
Last annotation update)

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QANEAT 1D
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QANEAT 1D
QANEAT 01-0
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Matches 101; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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InterPro; IPRO03990; Pancreatis_ac.
Pfam; PF00059; Lectin_C; 1.
PRINTS; PR01504; PNCERATITSAP.
SMART; SM00034; CLECT; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Namikawa K., Murakami K., Fukushima M., Kiyama H.; "Differential regulation of Reg family member expression periperal nerve injury."; Submitted (FEB-2004) to the EMBL/GenBank/DDBJ databases. EMBL; AB164049; BAD38673.1; -.
                                                                                                                                                                                                                                                                                                                                                    EMBL; AY126671; AAM95599.1; -.
HSSP; P21963; JJZN.
HSSP; P20005529; F:sugar binding; IE
InterPro; IPR001304; Lectin C.
InterPro; IPR003990; Pancreatis ac.
Pfam; PP00059; Lectin C; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-OCT-2002 (TrEMBLrel.
01-OCT-2002 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                  PRINTS; PR01504; PNCREATITSAP.
SMART; SM00034; CLECT; 1.
PROSSITE; PS50041; C. TYPE LECTIN 2; 1.
SEQUENCE 113 AA; 12832 MW; AZE9DF1A729C78DA CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kamarainen M., Heiskala K., Submitted (JUN-2002) to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REG-like protein splice
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q8NER7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q8NER7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN=Wistar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rattus norvegicus (Rat).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Regenerating islet-derived family member
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        105;
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157 AA; 18269 MW; 55B129FB10BA4D1D CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAWYLYRSWSG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MASRSMRLLLLLLSCLAKTGVLGDIIMRPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQS
                                     MASRSMRLLLLLLSCLAKTGVLGDIIMRPSCAPGWFYHKSNCYGYFRKLRUWSDAELECQS
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MASRSMRLLLLLSCLAKTGVLGDIIMRPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQS
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                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE=Ileum;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                        62.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   67.6%; Score 593.5; 66.5%; Pred. No. 2.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22, Created)
22, Last sequence update)
26, Last annotation update)
variant 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21; Mismatches
                                                                                                               1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Heiskala M., Andersson L.C.; EMBL/GenBank/DDBJ databases.
                                                                                                            Score 544; DB 2;
Pred. No. 1.2e-45;
1; Mismatches 0
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Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31;
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                                                                                                                                                                          Length
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; Murinae; Rattus.
                                                                                                                                                                             113;
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RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan R., Farmer A.A., Rubin G.M., Hong L.,
RA Hopkins R.F., Jordan R., Farmer A.A., Rubin G.M., Hong L.,
RA Hopkins R.F., Jordan R., Fonaldo M.F., Casavant T.L., Scheetz T.B.,
RA Hopkins R.F., Jordan R., Fonaldo M.F., Casavant T.L., Scheetz T.B.,
RA S.S., Loquellano N.A., Peters G.J., Abramson R.D., Millahy S.J.,
RA Raha S.S., McEwan P.J., McKernan K.J., Malek J.A., Gunaratine P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Hopkins M. M., Touchman J.W., Green B.D., Dickson M.C.,
Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Kzzywinski M.I., Skalska U., Smailus D.B., Schmerch A., Schein J.E.,
Van Genes S.J., Marra M.A.;
Query Match
Best Local S
Matches 75
                                                                                                                                           PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q7SZ75
Q7SZ75;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
Xenopodinae; Xenopus.
                                                                                                                                                                                                 GO; GO:0005529; F:sugar binding;
InterPro; IPRO1304; Lectin_C.
Pfam; PF00059; Lectin_C; 1.
SMART; SM00034; CLECT; 1.
                                                                                                                                                                                                                                                                                                                    Klein S., Strausberg R.;
Submitted (JUN-2003) to the
EMBL; BC053817; AAH53817.1;
HSSP; P22897; 1EGG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MGC64513 protein.
Xenopus laevis (African clawed frog).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-OCT-2003
01-OCT-2003
                                                                                                                  SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=Who
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               initiative.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Genetic and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Richardson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WEDLINE-22341132; PubMed=12454917; DOI=10.1002/dvdy.10174; WEDLINE-22341132; PubMed=12454917; DOI=10.1002/dvdy.10174;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   σ
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Similarity
                                                                                                               M00034; CLECT; 1.

PS00615; C_TYPE_LECTIN_1; 1.

PS50041; C_TYPE_LECTIN_2; 1.

160 AA; 18330 MW; 6A5502F24689179A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 225:384-391 (2002) .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Acad.
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genomic
                           49.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tools
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26,
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   35;
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, Last sequence up
, Last annotation :
                                                                                                                                                                                                                                                                                                                                                                                EMBL/GenBank/DDBJ databases.
Score 438.5; DB 2;
Pred. No. 4.1e-35;
5; Mismatches 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99:16899-16903 (2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Xenopus
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      Indels
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RESULT 7
Q8NER6
ID Q8NE
AC Q8NE
AC Q8NE
DT 01-C
DT 01-C
DT 01-C
DE REG-
OS Homc
OC Euka
OC Mamm
OC NCBI
RN 111
RP SEQI
RN 121
RP SEQI
RN 5EMBH
DR EMBH
SQ SEQI
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Best Local S
Matches 62
                                                    "Ostrich (Struthio camelus) eggshell matrix con
type lectin-like proteins Isolation, amino aci-
posttranslational modifications.";
Biochim. Biophys. Acta 1696:41-50(2004).
-!- SUBCELLULAR LOCATION: Eggshell matrix.
-!- SUBCELLULAR LOCATION: Eggshell matrix.
-!- MASS SPECTROMETRY: MW=15343.2; MW_ERR=4; ME
PANGE=1-132; NOTE=Ref.1.
-!- SIMILARITY: Contains 1 C-type lectin family
InterPro; IPR002353; AntifreezeII.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            QBNER6;
01-OCT-2002 (TrEMBLrel. 22, Created)
01-OCT-2002 (TrEMBLrel. 22, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
nEG-like protein splice variant 2.
Homo sapiens (Human).
                                                                                                                                                                                                                                   SEQUENCE, SUBCELLULAR LOCATION, AND MASS SPECTROME
TISSUE-Eggshell matrix;
PubMed=14756203; DOI=10.1016/j.bbapap.2003.09.006;
Mann K., Siedler F.;
                                                                                                                                                                                                                                                                                                                                                                                                               Struthiocalcin-1 (SCA-1). Struthio camelus (Ostrich).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kamarainen M., Heiskala K.,
Submitted (JUN-2002) to the
EMBL, AY126672; AAM95800-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUL-2004
05-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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                  InterPro; IPR001304; Lectin_C.
InterPro; IPR003990; Pancreatis_ac.
                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota;
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                                                                                                                                                                                                                                                                                                                              CBI_TaxID=8801;
                                                                                                                                                                                                                                                                                                                                                                         rchosauria; Aves;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRCA
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                                                                                                                                                                                                                                                                                                                                                                                             Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   134 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                             (Rel. 44, Created)
(Rel. 44, Last sequence update)
(Rel. 44, Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                         ; Chordata; Craniata; Vertebrata; Palaeognathae; Struthioniformee;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14993 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 2.60
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 307.5;
Pred. No. 2.6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
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                                                                                                                                                                                                                                                                                              AND MASS SPECTROMETRY
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                                                                                                                                                                                             matrix contains two amino acid sequence
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                                                                            family domain
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                                                                                                                  METHOD=Electrospray;
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Struthionidae;
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e, and
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ACAL_AN
ACAL_AN
      TINGE BERREER RECORDED TO THE PROPERTY OF THE 
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Best Local S
Matches 50
                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUB-ENgushell matrix;

TISSUB-ENgushell matrix;

MEDLINE-22439773; PubMed=12431998; DOI=10.1074/jbc.M201518200;

Lakehminarayanan R., Valiyaveettil S., Rao V.S., Kini R.M.;

"Purification, characterization, and in vitro mineralization sof a novel goose eggshell matrix protein, ansocalcin.";

of a novel goose eggshell matrix protein, ansocalcin.";

J. Biol. Chem. 278:3928-2936(2003).
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DISULFID
DISULFID
SEQUENCE
PROSITE; PS00615; C_TYPE_LECTIN_1; 1.
PROSITE; PS50041; C_TYPE_LECTIN_2; 1.
Direct protein sequencing; Lectin.
DOMAIN
1 132 C-type lectin.
DISULFID 3 14 By similarity.
                                                                                              Pfam, PF00059; Lectin C; 1.
PRINTS; PR00356; ANTIFREEZEII.
PRINTS; PR01504; PNCREATITSAP.
SMART; SM00034; CLECT; 1.
                                                                                                                                                                                                                                                                                                                -!- SUBUNIT: Homodimer or homotrimer.
-!- SUBCELLULAR LOCATION: Eggshell matrix.
-glands on the walls of oviduct and inco-
structure during its formation.
                                                                                                                                                                                                                                                                                                                                                                                                     J. Biol. Chem. 278:2928-2936(2003).
-!- FUNCTION: Induces spherical aggregates of vitro. Believed to play an active role in
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PRINTS; PR01504; PNCREATITSAP.
SMART; SM00034; CLECT; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SPECTROMETRY
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05-JUL-2004
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28-FEB-2003
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DOMAIN 10
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                                                                                                                                                                                                                                       HSSP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=8844;
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                                                                                                                                                                                                                 InterPro;
                                                                                                                                                                                                                                                        -!- SIMILARITY: Contains 1 C-type
                                                                                                                                                                                                                                                                                              -!- MASS SPECTROMETRY: MW=15342; METHOD=Electrospray; RANGE=1-132
                                                                                                                                                                             InterPro;
                                                                                                                                                                                               InterPro;
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                                                                                                                                                                                                                                                                                                                                                                                             calcification.
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                                                                                                                                                                         IPR002353; AntifreezeII.
IPR001304; Lectin_C.
IPR003990; Pancreatis_ac.
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0 129 C-type lectin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SUBUNIT,
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Pred. No. 1.1e
20; Mismatches
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                                                                                                                                                                                                                                                                                                                                rix. May be present incorporated into
                                                                                                                                                                                                                                                        family domain
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MASS

Euteleostomi;

Anser

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30 CAPGWFYHKSNCYGYFRKLRNWSDAELECQSYGNGAHLASILSLKEASTIAEYISGY---
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                                                                                                                                                                                                                                                                                                                          CPKGWLDFRGNCYGYFRYELPWKRAEAWCRSIRAGAHLASIHTSEEHRAIAKFISQYHHG
41, Last sequence update)
44, Last annotation update)
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By similarity.
By similarity.
By similarity.
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LITIMOUSE
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                                                                                                             RP SEQUENCE FROM N.A.

RP STRAIN=CS7BL/6J; TISSUE=Thymus;

RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.24260389;

RX MEDLINE=22388257; DoI=10.1073/pnas.24260389;

RX MEDLINE=22388257;

RX MEDLINE=2238825;

RX MEDLINE=223
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P43137;
01-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DISULFID
DISULFID
SEQUENCE
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05-JUL-2004 (Rel. 44, Last annotation update)
Lithostathine 1 precursor (Pancreatic stone protein 1) (PSP)
(Pancreatic thread protein 1) (PTP) (Islet of langerhans regi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Moriizumi S., Okamoto H., Nakagawara K.-I., Watanabe T., Miyashita "Structure, chromosomal localization, and expression of mouse r genes, reg I and reg II. A novel type of reg gene, red IT ....' the mouse genome.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=C57BL/6J;
MEDLINE=93340209; PubMed=8340418;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
                                                                                                   Proc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Name=Reg1;
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      FUNCTION: Might act as an carbonate precipitation.
TISSUE SPECIFICITY: Expres
                                                                                                Natl. Acad.
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103
132
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                                                                                                Sci. U.S.A.
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      Expressed only in
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Pred. No. 1.
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                                                                                                99:16899-16903 (2002)
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y similarity.
36CE42EA4572E6B7 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         165
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regenerating islets and
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                                                                spontaneous calcium
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RESULT 11
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Best Local S
Matches 52
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InterPro; IPR002353; AntifreezeII.
InterPro; IPR001304; Lectin_C.
InterPro; IPR001390; Pancreatis_ac.
Pfam; PP00059; Lectin_C; 1.
PRINTS; PR00356; ANTIFREEZEII.
PRINTS; PR00356; ANTIFREEZEII.
SMART; SM00034; CLECT; 1.
                                                                                                     01-JUL-1989 (Rel. 11, Created)
01-JUL-1989 (Rel. 11, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation update)
Lithostathine precursor (Pancreatic stone protein) (PSP) (Pancreatic thread protein) (PTP) (Islet of langerhans regenerating protein) (REC (Islet cells regeneration factor) (ICRF).
                                                                                                                                                                                    LITH RAT
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DISULFID
DISULFID
CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           the European Bioinformatics Institute. There are no rest
use by non-profit institutions as long as its content
modified and this statement is not removed. Usage by an
entitles requires a license agreement (See http://www.isb
or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HSSP;
           SEQUENCE FROM N.A. MEDLINE=91093273; PubMed=1985964;
                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                Name=Reg1; Synonyms=Reg;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Glycoprotein;
SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PROSITE; PS00615; C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; D14010;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN
                                            NCBI_TaxID=10116;
                                                                                     Rattus norvegicus (Rat).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PIR; A47148; A47148.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SIMILARITY: Contains 1 C-type lectin family domain.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  normal exocrine pancreas, but not in normal pancreatic islets. 
Expressed strongly in pancreas, moderately in gallbladder, and
                                                                                                                                                                                                  RAT
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                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
52; Conser
s.
                                                                                                                                                                                                                                                                            YLYRSW-SGKSMGGNK-HCAEMSSNNNFLTWSSNECNKRQHFLCKYR
                                                                                                                                                                                                                                                                                                            ELECQSYGNGAHLASILSLKEASTIAEYI--SGYQRSQPIWIGLHDPQKRQQWQWIDGAM 112
                                                                                                                                                                                                                                                               FLYKSWATGSPNSSNRGYCVSLTSNTGYKKWKDDNCDAQYSFVCKFK
                                                                                                                                                                                                                                                                                                                                                            ILLSCL-----IVLSPSQGQEAEEDLPSARISCPEGSNAYSSYCYYFTEDRLTWADA
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                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
 Verdier J.M.,
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161
153
129
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165
163
                                                             Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE_LECTIN_1;
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Lithostathine 1.
C-type lectin.
By similarity.
By similarity.
By similarity.
By similarity.
By similarity.
N-linked (GlCNNG...
W; 2950174AF5D666BA (
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 254.5;
Pred. No. 5.2
                                                             Craniata; Veri
Sciurognathi;
                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                  165
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 Dagorn J.-C., Giorgi D.;
                                                                                                                                                                                                                                                                                                                                                                                                                          5.2e-17;
                                                             Vertebrata;
thi; Muridae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                             51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              CRC64
                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 165;
                                                             Euteleostomi;
Murinae; Rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Potential)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             restrictions
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                                                              Rattus
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Query Match
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                                       DISULFID
CARBOHYD
SEQUENCE
                                                                              DOMAIN
DISULFID
DISULFID
                                                                                                                                    Direct
SIGNAL
                                                                                                                                                                                                                                                                              EMBL; L07512; AAA41533.1; --
EMBL; M62930; AAA41974.1; --
EMBL; M16962; AAA42028.1; --
EMBL; D26164; BAA05149.1; --
EMBL; D26164; BAA05149.1; --
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-90031455; PubMed=2680252;
Adrich Z., de Caro A.M., Guidoni A.A., Woudstra M.E., Rovery M.
Characterization in rat pancreatic juice of a protein homologo
the human pancreatic stone protein.";
Comp. Biochem. Physiol. 93B:793-797(1989).
-I- FUNCTION: Might act as an inhibitor of spontaneous calcium
carbonate precipitation.
-I- TISSUE SPECIFICITY: Expressed only in regenerating islets,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Rapid
gene.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Rat pancreatic stone protein messenger RNA. Abundant expression in mature exocrine cells, regulation by food content, and sequence identity with the endocrine reg transcript.";
J. Biol. Chem. 266:786-791/1001\
                                                                                                                      CHAIN
                                                                                                                                                                                                                                                                                                                                                                            modified and this statement is not removed. entities requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Enthe European Bioinformatics Institute. There are no restruse by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dusetti N.J., Frigerio "Rapid PCR cloning and
                                                                                                                                                                                                                                                                                                                                                              entitles requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Miyashita H., Suzuki Y., Watanak
YoneKura H., Okamoto H.;
"Structure and characterization
Seikagaku 65:1082-1082(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=93326645; PubMed=7916640; DOI=10.1016/0167-4781(93)90100-R; Dusetti N.J., Frigerio J.M., Dagorn J.-C., Iovanna J.L.;
                                                                                                                                            PROSITE; PS00615; C_TYPE_LECTIN_1; 1.
PROSITE; PS50041; C_TYPE_LECTIN_2; 1.
Direct protein sequencing; Glycoprotein; Lectin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in normal pancreatic islets, insulinomas or regenerating -!- SIMILARITY: Contains 1 C-type lectin family domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Terazono K., Yamamoto H.,
Tochino Y., Okamoto H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-88115343; PubMed-2963000; Terazono K., Yamamoto H., Takasawa
                                                                                                                                                                                                 PRINTS;
                                                                                                                                                                                                               InterPro; IPRO01304; Lectin_C.
InterPro; IPR003990; Pancreatis_ac.
Pfam; PF00059; Lectin_C; 1.
                                                                                                                                                                                                                                                       RGD; 3552; Reg1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FISSUE-Pancreas;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN-Wistar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biochim. Biophys. Acta 1174:99-102(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "A novel gene activated in regenerating J. Biol. Chem. 263:2111-2114(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE OF
                                                                                                                                                                                                                                                                    P05451; 1LIT.
                                                                                                                                                                                     ; PR01504; PNCREATITSAP.
SM00034; CLECT; 1.
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                                                     163
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161
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129
                                        18672 MW;
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                                                  Lithostathine.
C-type lectin.
By similarity.
By similarity.
By similarity.
By similarity.
N-linked (GlcNAc.
Score
Pred.
                                        9B61EB236B82CF8A
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254.5; DB 1;
No. 5.2e-17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rat Reg I
                                                                                                                                                                                                                                                                                                                                                                            (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shiga K., Yonemura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     islets.";
                                                                                                                                                                                                                                                                                                                                                                                     Usage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     of the rat lithostathine
                                         CRC64;
                                                                                                                                              Signal.
            Length
                                                   (Potential)
                                                                                                                                                                                                                                                                                                                                                                                                                 he EMBL outst
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                                                                                                                                                                                                                                                         Direct
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE-Eggshell matrix;
PubMed=14726203; DOI=10.1016/j.bbapap.2003.09.006;
Mann K., Siedler F.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Struthiocalcin-2 (SCA-2). Struthio camelus (Ostrich)
                                                                                                                                                                                                                                                                 PROSITE; PS00615; C_TYPE_LECTIN_1; PROSITE; PS50041; C_TYPE_LECTIN_2; Direct protein sequencing; Lectin;
                                                                                                                                                                                                                                                                                                     PRINTS; PR01504; PNCREATITSAP. SMART; SM00034; CLECT; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-JUL-2004
05-JUL-2004
                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                    8
                                                                                                                                                                                                           8
                                                                                                                                                                                                                                                                                                                                                           RANGE=1-142; NOTE=Ref.1.
                                                                                                                                                                                                                                                                                                                                                                                  -!- SUBCELLULAR LOCATION: Eggshell matrix.-!- MASS SPECTROMETRY: MW=16834.1; MW_ERR=2; METHOD=Electrospray;
                                                                                                                                                                                                                                                                                                                                                                                                           Biochim. Віорһув. Acta 1696:41-50(2004).
                                                                                                                                                                                                                                                                                                                                                                                                                                 "Ostrich (Struthio camelus) eggshell matrix type lectin-like proteins. Isolation, amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AND SER-68,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=8801;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Archosauria; Aves;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-JUL-2004
                                                                                                                                                                                                                                                                                                                                   InterPro; IPR003990; Pancreatis_ac.
                                                                                                                                                                                                                                                                                                                                                                                                                      posttranslational modifications.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE,
                                                                                                                                                                                                ĝ
                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR001304; Lectin_C.
                                                                                                                                                                                    RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRCA
            139
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                                                                                                                                                                                                                                                                                                                             PF00059; lectin
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                                 63
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                                                                                w
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                                                                                                                                       Similarity
LTWSSNECNKRQHFLCKYR
                                                                                                     RPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQSYGNGAHLASILSLKEASTIAEYISGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GSLFLYKSWDTGYPNNSNRGYCVSVTSNSGYKKWRDNSCDAQLSFVCKFK 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GAMYLYRSW-SGKSMGGNK-HCAEMSSNNNFLTWSSNECNKRQHFLCKYR
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                                                                             RAGCAKGWIPFDGRCYGFFPQELSWRRAEGFCQRLGARTHLASIHSEEEHQAIVSWLASS
                                   OPYSDSEEEAGEEVWIGLHRPLGRRNWEWSDGTKLDYGSWYRDVFLRRRACVALEDTTDF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AEADLFCQNMNSG-YLVSVLSQAEGNFLASLIKESGTTAAN-VWIGLHDPKNNRRWHWSS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBCELLULAR LOCATION, PHOSPHORYLATION SITES B, AND MASS SPECTROMETRY.
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(Rel.
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44, Last sequence update)
44, Last annotation update)
                                                       RSQPIWIGLHDPQKRQQWQWIDGAMYLYRSWSGKSMGGNKHCAEMSSNNNF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Palaeognathae;
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By similarity.
By similarity.
By similarity.
Phosphoserine.
Phosphoserine.
                                                                                                                                      Score 247; DB 1;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                       C-type lectin
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                                                                                                                            66;
                                                                                                                                                                         CRC64;
                                                                                                                                                  Length 142
                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence,
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                                                                                                                            Indels
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Struthionidae;
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01-NOV-1997 (Rel. 3
01-NOV-1997 (Rel. 3
05-JUL-2004 (Rel. 4
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DISULFID
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                          PRINTS; PRUIDOT, I. SMART; SMO0034; CLECT; 1.

PROSITE; PS00615; C_TYPE_LECTIN_1; FALSE_NEG.

PROSITE; PS50041; C_TYPE_LECTIN_2; 1.

PROSITE; PS50041; C_TYPE_LECTIN_2; 1.

Acute phase; Inflammatory response; Lectin; Multigene family; Signal.

Potential.

Potential.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; D63361; BAA18930.1
EMBL; D63362; BAA18931.1
HSSP; P05451; 1LIT.
MGD; MGI:109406; Reg3g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR001304; Lectin C.
InterPro; IPR003990; Pancreatis ac.
Pfam; PF00059; Lectin C; 1.
PRINTS; PR01504; PNCREATITSAP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   encoding type III Reg, RegIII alpha, Gene 185:159-168(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Name=Pap3; Synonyms=Reg3g;
Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JUL-2004 (Rel. 44, Last annotation update)
Pancreatitis-associated protein 3 precursor (REG III-gamma)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PAP3 MOUSE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Structure, chromosomal localization and expression of mouse genes encoding type III Reg, RegIII alpha, RegIII beta, RegIII gamma.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               healthy pancreas.
SIMILARITY: Contains 1 C-type lectin family domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FUNCTION: Might be a stress protein involved in the control of bacterial proliferation.
SUBCELLULAR IOCATION: Secreted.
TISSUB SPECIFICITY: Constitutively expressed in the small
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       intestine, moderately in colon and
                                                                                                  13
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73
                                             63
                                                                                                                                                                                                                              Similarity
SG-HLVSVLSGAEASFLSSMIKSSGNSGQYVWIGLHDPTLGYEPNRGGWEWSNADVMNYI
                                          NGAHLASILSLKEASTIAEYI-SGYQRSQPIWIGLHDP-----QKRQQWQWIDGAMYLYR
                                                                                               MILISCLMLLSQVQGEVAKKDAPSSRSSCPKGSRAYGSYCYALFSVSKNWYDADMACQKRP
                                                                                                                                               LLLSCL-AKTGVLGDII-----MRPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQSYG
                                                                                                                                                                                                                                                                                                      1
27
38
40
68
145
174 AA;
                                                                                                                                                                                                                                                                                                                                                   ; Inflammatory
1 26
27 174
38 174
40 51
68 170
                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                        19307 MW;
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C-type lectin.
By similarity.
By similarity.
By similarity.
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Pred. No. 3.4e-16;
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                                                                                                                                                                                                    Mismatches
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RESULT 14
Q6TRS6
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Best Local S
Matches 48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A Kassab B.H., de Carvalho D.D., Ollveira M.A., Baptista G.R.,
A Pereira G.A., Novello J.C.;
T "Cloning, expression, and structural analysis of recombinant BJC
T c-type lectin from the Bothrops jararacussu snake venom.";
Protein Expr. Purif. 35:344-352(2004).
L Protein Expr. Purif. 35:344-352(2004).
R EMBL; AY38642; AAQ92957.1; --
R GG; GG:0005529; F:sugar binding; IEA.
R InterPro; IPR001304; Lectin C.
R InterPro; IPR003990; Pancreatis_ac.
R Ffan; PF00059; Lectin C; 1.
R PRINTS; PR01504; PNCREATITSAP.
R SMART; SM00034; CLECT; 1.
                                                                                                                                                                                                                     Q6QX33; PRELIMINARY,
Q6QX33;
05-JUL-2004 (TrEMBLrel. 2
05-JUL-2004 (TrEMBLrel. 2
05-JUL-2004 (TrEMBLrel. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi; Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea; Viperidae; Crotalinae; Bothrops.
NCBI_TaxID=8726;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JUL-2004
05-JUL-2004
                                                                 NCBI_TaxID=8723;
                                                                                                Viperidae; Crotalinae;
                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata;
                                                                                                                                                                         Bothrops insularis (Island
                                                                                                                                                                                                C-type lectin.
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PROSITE; PS50041; C TYPE LECTIN 2; 1.
CHAIN 24 157 BJcul.
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Kassab B.H., de Carvalho D.D., Oliveira M.A., B
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TISSUE=Venom gland
                          SEQUENCE FROM N.A.
                                                                                                                       Lepidosauria; Squamata; Scleroglossa;
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llarity 36.9%;
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ta; Craniata; Vertebrata; Eute
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Pred. No. 4.5e-16;
6; Mismatches 62
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                                                                                                                               Colubroidea;
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Search completed: February 11, 2005, 23:32:39 Job time : 179 secs
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Junqueira-de-Azevedo I.L.M., Dutra D.L.S., Pujol-Luz M., Castro H.C.,

A Ho P.L., Zingali R.B.;

EL Submitted (JAN-2004) to the EMBL/GenBank/DDBJ databases.

EMBL; AY522720; AAS01426.1; -

GO; GO:0005529; F:sugar binding; IEA.

InterPro; IPR002353; AntifreezeII.

InterPro; IPR001304; Lectin C.

InterPro; IPR001304; Lectin C.

InterPro; IPR001304; Lectin C.

R InterPro; IPR001304; Lectin C.

R PRINTS; PR0059; Lectin C; 1.

R PRINTS; PR00356; ANTIFREEZEII.

R PRINTS; PR00356; ANTIFREEZEII.

R PRINTS; PR00356; ANTIFREEZEII.

R PROSITE; PS00615; C_TYPE_LECTIN_1; 1.

PROSITE; PS00615; C_TYPE_LECTIN_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 27.7%; Score 243.5; DB 2; Length 158; Best Local Similarity 34.0%; Pred. No. 6e-16; Matches 51; Conservative 22; Mismatches 68; Indels 9.
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SEQUENCE
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                                                                                                                                                                                                        126 KEFCVELVSDTGYRLWNDQVCESKNAFLCQ 155
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Result
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Maximum Match 100%
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Perfect score:
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       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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length: 2000000000
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878
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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     GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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     DB
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regenerating islet
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pancreatitis assoc
agkisacutacin alpha
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reg II, regenerati
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                                                                                        brevican - human
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RESULT 2 A28351 pancreati N;Alternat C;Species C;Date: 3 C;Accessi R;Terazon J. Biol. (A;Title: / A;Referen A;Recessi A;Recessi A;Residue	RESULT 1 A47148 reg I, regenerating C;Species: Mus muscul C;Date: 03-May-1994 C;Accession: A47148 R;Unno, M.; Yonekura, J. Biol. Chem. 268, A;Title: Structure, A;Reference number: A A;Residues: preliminary A;Molecule type: DNA A;Residues: 1-165 < ON A;Cross-references: A;Introns: 2/1/1; 60/2 C;Superfamily: tetran R;35-161/Domain: C-ty P;35-46,63-161,136-11 Query Match Best Local Similari Matches 52; Cons Matches 52; Cons Oy 10 LLLSCLJ Oy 11 SLECQS: Oy 13 YLYRSW Oy 11 STYRSW Db 118 FLYKSW		4444444 544444 54444 5444 5444 5444 54
RESULT 2 A28351 pancreatic stone protein precursor - rat N;Alternate names: lithostathine C;Species: Rattus norvegicus (Norway rat) C;Date: 31-Mar-1990 #sequence_revision 31- C;Accession: A28351; A39081; FD10147; S3461 R;Terazono, K., Yamamoto, H.; Takasawa, S. J. Biol. Chem. 263, 2111-2114, 1988 A;Title: A novel gene activated in regener A;Reference number: A92704; MUID:88115343; A;Accession: A28351 A;Accession: A28351 A;Residues: 1-165 <ter></ter>	RESULT 1 A47148 reg I, regenerating islet cells - mouse C; Species: Mus musculus (house mouse) C; Date: 0.3-May-1994 #sequence_revision & A-Mayawara, K.; Wa C; Accession: A47148 R; Unno, M.; Yonekura, H.; Nakagawara, K.; Wa J. BiOl. Chem. 268, 15974-15982, 1993 A; Title: Structure, Chromopomal localization A; Reference number: A47148; MUID:93340209; P A; Accession: A47148 A; Residues: 1-165'CUNN' A; Roccise type: DNA A; Residues: 1-165'CUNN' A; Cross - references: UNIPROT: P43137; GB:D1401 C; Genetics: A711; 60/3; 106/3; 144/1 C; Genetics: C-type lectin homology <lch #sta="" 10="" 31.1%;="" 35-161'domain:="" 35-266,63-161,136-153="" 35;="" 52;="" 8="" <lch="" best="" bonds:="" c-type="" conservative="" db="" disulfide="" f;="" homology="" illscl<="" lectin="" lllsclaktgvlgdiimrp="" local="" match="" matches="" mismatc="" no.="" pred.="" query="" qy="" similarity="" td=""><td></td><td>185 21.1 883 185 21.1 1257 84.5 21.0 146 82.5 20.8 301 181 20.6 291 181 20.6 3562 119 20.4 2397 179 20.4 2409 20.4 2409 20.5 162 277.5 20.2 162 177 20.2 163 177 20.2 3381 177 20.2 3381 177 20.2 3381</td></lch>		185 21.1 883 185 21.1 1257 84.5 21.0 146 82.5 20.8 301 181 20.6 291 181 20.6 3562 119 20.4 2397 179 20.4 2409 20.4 2409 20.5 162 277.5 20.2 162 177 20.2 163 177 20.2 3381 177 20.2 3381 177 20.2 3381
Mar-1990 #tex ; Shiga, K.; ating islets. PMID:2963000	O9-Ma ation 09; Wa clCH #sta 254. MAEYI ASLIK ASLIK NNELT	ALIGNMENTS	2 S49126 2 S28764 2 JC7135 2 S13165 2 S13165 2 PC7027 1 LNHU1 2 R47171 2 S47171 2 S52781 1 A60979 1 A60979 1 LNRC1 1 LNRC1 1 LNRC3 2 T14274 2 T14274 2 T14274 2 T14274 2 T14274 2 T14274
t_change 09-Jul-2004 Yonemura, Y.; Tochino, Y.; Okamot	y-1994 #text_change 09-Jul-2004 tanabe, T.; Miyashita, H.; Moriizumi, S.; Ok , and expression of mouse reg genes, reg I a MID:8340418 0; NID:g391771; PIDN:BAA03111.1; PID:g391772 tus predicted tus predicted 5; DB 2; Length 165; 9.2e-18; 9.2e-18; 9.2e-18; 9.3e-18; 9.3e-18; 9.3e-18; 1		brevican precursor neurocan precursor agkisacutacin beta asialoglycoprotein aggretin alpha cha hepatic lectin H1 chondroitin sulfat neurocan - mouse versican precursor versican precursor asialoglycoprotein lectin BRA3-1 prec lectin BRA3-2 prec versican precursor versican precursor versican precursor

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A; Experimental source: egg-shell C; Superfamily: tetranectin; C-ty
                                                                                                                                                    C;Accession: S78596
R;Baernholdt, D.; Andersen,
submitted to the Protein Sec
                                                                                                                                                                                                                                                                                 C;Species: Gallus gallus (chicken)
C;Date: 24-Oct-1998 #sequence_revision
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S78596
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Biochim. Biophys. Acta 1174, 99-102, 1993
A;Title: Rapid PCR cloning and sequence determination of the rat lithostathine A;Reference number: S34618; MUID:93326645; PMID:7916640
A;Accession: S34618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Experimental source: pancreas
R;Dusetti, N.J.; Frigerio, J.M.; Dagorn, J.C.; Iovanna,
Biochim. Biophys. Acta 1174, 99-102, 1993
Biochim. Biophys. Acta 1174, 99-102, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: GB:M62930; GB:J05722; NID:g206462; PIDN:AAA41974.1; PID:g206463 R;Adrich, Z.; De Caro, A.M.; Guidoni, A.A.; Woudstra, M.E.; Rovery, M. Comp. Biochem. Physiol. B 93, 793-797, 1989 A;Title: Characterization in rat pancreatic juice of a protein homologous to the human A;Reference number: PL0147; MUID:90031455; PMID:2680252 A;Accession: PL0147
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                                                                                                                          A; Molecule type: protein A; Residues: 1-142 < BAE>
                                                                                                                                                                                                                                                                                                                                         ovocleidin -
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C; Superfamily: tetranectin; C-type lectin homology
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A; Residues: 1-165 < DUS>
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A; Residues: 1-165 < ROU>
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A;Accession: A39081
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;22-165/Product: pancreatic stone protein #status predicted <MAT>;35-161/Domain: C-type lectin homology <LCH>;26/Modified site: pyrrolidone carboxylic acid (Gln) (in mature form)
       ;61,67/Binding
                                                     ;Keywords:
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Best Local
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rds: phosphoprotein

33-138,113-130/Disulfide bonds:

,33-138,113-170/Disulfide bonds:
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les 52; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GAMYLYRSW-SGKSMGGNK-HCAEMSSNNNFLTWSSNECNKRQHFLCKYR 157
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                                                                      C-type lectin homology
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       (covalent) #status ex
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    #status experimental
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A;Gene: GDB:REG1B; REG1
A;Cross-references: GJB:342079
A;Cross-references: GJB:342079
A;Map position: 2p12-2p12
A;Introns: 22/1; 61/3; 107/3; 145/1
A;Introns: 22/1; 61/3; 107/3; 145/1
C;Superfamily: tetranectin; C-type lectin homology
C;Reywords: glycoprotein; lectin; panoreas; pyroglutamic acid
F;1-22/Domain: signal sequence #status predicted <SIG>F;1-22/Domain: signal sequence #status predicted <SIG>F;3-166/Product: regenerating islet lectin lbeta #status predicted
F;34-166/Product: pancreatic stone protein #status predicted cMAT2>
F;36-162/Domain: C-type lectin homology <CCH>A
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A;Molecule type: DNA
A;Residues: 1-166 <MO2>
A;Residues: 1-166 <MO2>
A;Cross-references: GB:D17291; NID:g474307; PIDN:BAA04124.1; PID:g474308
A;Cross-references: GB:D17291; NID:g474307; PIDN:BAA04124.1; PID:g474308
A;Cross-references: GB:D17291; NID:g474307; PIDN:BAA04124.1; PID:g474308
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A;Residues: 1-166 <BAR>
A;Residues: 1-166 <BAR>
A;Residues: 1-166 <BAR>
A;Residues: UNIPROT:P48304; GB:L08010; NID:g307368; PIDN:AAA18204.1; PID:g487726
A;Cross-references: UNIPROT:P48304; GB:L08010; NID:g307368; PIDN:AAA18204.1; PID:g487726
A;Cross-references: UNIPROT:P48304; GB:L08010; NID:g307368; PIDN:AAA18204.1; PID:g487726
A;Moriizumi, S.; Watanabe, T.; Unno, M.; Nakagawara, K.; Suzuki, Y.; Miyashita, H.; Yone'
Biochim. Biophys. Acta 1217, 199-202, 1994
A;Title: Isolation, structural determination and expression of a novel reg gene, human r.
A;Reference number: S42729; MUID:94153997; PMID:8110835
A;Accession: S42729
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RGHU1B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence revision 03-Aug-1995 #text_change
C;Accession: S34591, S42729; A44712
R;Bartoli, C.; Gharib, B.; Giorgi, D.; Sansonetti, A.; Dagorn, J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          regenerating islet lectin 1-beta precursor - human N;Alternate names: reg-related protein; reg1-beta protein N;Contains: pancreatic stone protein (PSP)
                                                                                                                                                                                                                                                            F,36-47,64-162,137-154/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                                                                      F;23/Modified site: pyrrolidone carboxylic acid (Gln) (in mature F;27/Binding site: carbohydrate (Thr) (covalent) #status predicte F;33-34/Cleavage site: Arg-Ile (trypsin) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: GB:D16816; NID:g474305; PIDN:BAA04091.1; A;Accession: A44712
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30 CAPGWFYHKSNCYGYFRKLRNWSDAELECQSYGNGAHLASILSLKEASTIAEYI-----
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RISCPEGTNAYRSYCYYFNEDPETWVDADLYCQNMNSG-NLVSVLTQAEGAFVASLIKES
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                                                               RPSCAPGWFYHKSNCYGYFRKLRNWSDAELECOSYGNGAHLASILSLKEASTIAEYISGY 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       289-293, 1993
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                                                                                                                                                             Score 240; DB 1;
Pred. No. 2.5e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 241.5; DB : Pred. No. 1.5e-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 D.; Sansonetti, A.; Dagorn, J.C.; Berge-Lefranc,
                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                            DB 1; Length 166
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R;Suzuki, Y.; Yonekura, H.; Watanabe, T.; Unno, M.; Moriizumi, Gene 144, 315-316, 1994
A;Title: Structure and expression of a novel rat RegIII gene.
A;Reference number: I60296; MUID:94314238; PMID:8039722
A;Accession: I60296
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    regenerating protein III (reg III) - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 29-May-1998 #sequence_revision 29-May-1998 #text_change 09-Jul-2004
C;Accession: I60296; I83377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R;Unno, M., Yonekura, H.; Nakagawara, K.; Watanabe, T.; Miyashit J. Biol. Chem. 268, 15974-15982, 1993
A;Title: Structure, Chromosomal localization, and expression of A;Reference number: A47148; MUID:93340209; PMID:8340418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         reg II, regenerating islet cells - mouse
C;Species: Mus musculus (house mouse)
C;Date: 03-May-1994 #sequence_revision 03-May-1994
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183377
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A; Residues: 1-173 < UNN>
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A;Status: preliminar
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                                                                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-174 < RE2>
                                                                                                                                                                                                                                                                                                   A; Molecule type: mRNA
A; Residues: 1-174 < RES>
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Best Local S
Matches 50
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;43-169/Domain: C-type lectin homology <LCH>
;43-54,71-169,144-161/Disulfide bonds: #status predicted
                                                                                                                                                                                                                                  Status: preliminary; translated
                                                                                                                                                                                                                                                                                                                                           Status: preliminary; translated from GB/EMBL/DDBJ
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                                                            ;Introns: 25/1; 64/3; 110/3; 153/1
;Superfamily: tetranectin; C-type lectin homology
;39-170/Domain: C-type lectin homology <LCH>
                                                                                                                           Gene: RegIII
                                                                                                                                                                        Cross-references: GB:D26078; NID:g471159; PIDN:BAA05071.1;
                                                                                                                                                                                                                                                                          Cross-references:
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26.9%; Score 236.5; DB 2 ilarity 28.7%; Pred. No. 5.8e-16; Conservative 33; Mismatches 72
                                                                                                                                                                                                                                                                               UNIPROT: P35231; GB: D23676; NID: g471157;
26.9%;
35.3%;
Score 236.5; DB 2;
Pred. No. 5.8e-16;
                                                                                                                                                                                                                                  from GB/EMBL/DDBJ
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                    Length
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                                                                                                                                                                        PID:g471160
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lectin, galactose-specific - western diamondback rattlesnake (;Species: Crotalus atrox (western diamondback rattlesnake) C;Date: 23-Aug-1991 #sequence_revision 23-Aug-1991 #text_change 09-Jul-2004 C;Accession: A38609
R;Hirabayashi, J.; Kusunoki, T.; Kasai, K.
J. Biol. Chem. 266, 2320-2326, 1991
A;Title: Complete primary structure of a galactose-specific lectin from the A;Reference number: A38609; MUID:91115849; PMID:1989986
A;Accession: A38609
                                     A;Cross-references: UNIPROT:P42854; EMBL:U09193; NID:g483931; PIDN:AAA79231.1; R;Frigerio, J.M.; Dusetti, N.J.; Garrido, P.; Dagorn, J.C.; Iovanna, J.L. Biochim. Biophys. Acta 1216, 329-331, 1993
A;Title: The pancreatitis associated protein III (PAP III), a new member of the A;Reference number: S43438; MUID:94060113; PMID:8241280
                                                                                                                                                                                                                               A;Title: Cloning, expression and chromosomal localization A;Reference number: 854979; MUID:95234061; PMID:7717998 A;Accession: 854979
A;Status: preliminary A;Molecule type: DNA A;Residues: 1-174 <DUS A;Resi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Dusetti, N.J.; Frigerio, Biochem. J. 307, 9-16, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pancreatitis-associated protein PAP-3 - rat
C;Species: Rattus norvegicus (Norway rat)
C;Date: 23-Aug-1995 #sequence revision 19-Oct-1995 #text_change 09-Jul-2004
C;Accession: S54979; S43438
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S54979
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A;Molecule type: protein
A;Residuse: 1-135 AIRs
A;Cross-references: UNIPROT:P21963
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A38609
A; Accession: S43438
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A;Molecule type: protein
A;Residues: 38-138;141-175 <CAI>
A;Residues: 38-138;141-175 <CAI>
C;Comment: The purified protein undergoes a reversible globule-fibril transformation C;Superfamily: tetranectin; C-type lectin homology
C;Keywords: disulfide bond; extracellular protein; pancreas
C;Keyroduct: pancreatic thread protein chain A #status experimental <ACH>
F;38-138;Product: pancreatic thread protein chain B #status experimental <BCH>
F;141-175/Product: pancreatic thread protein chain B #status experimental <BCH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Cross-references: UNIPROT:P23132; GB:M59794; NID:g163648; PIDN R;Cai, L.; Harris, W.R.; Marshak, D.R.; Gross, J.; Crabb, J.W. J. Protein Chem. 9, 623-632, 1990
A;Title: Structural analysis of bovine pancreatic thread protein A;Reference number: A53897; MUID:91197388; PMID:2085387
A;Accession: A53897
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pancreatitis-associated C; Species: Rattus norve
                                     A48689
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A; Residues: 1-175 < DEL>
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C;Genetics:
                                                             RESULT 10
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                                                                                                                       NYVAWETDPAAISSPGYCGSLSRSSGYLKWRDHNCNLNLPYVCKF
                                                                                                                                                                                                   KRPSG-HLVSVLSGAEESFVASLVRNNLNTQSDIWIGLHDPTEGSEANAGGWEWISNDVL
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30.9%; Pred. No. 2e-15;
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C;Species: Homo sapiens (man)
C;Date: 31-Mar-1990 #sequence revision 03-Aug-1995 #text change 09-Jul-2004
C;Accession: A35197; B28351; S12950; S02767; S02419; S00713; S01471; A25246
R;Watanabe, T.; Yonekura, H.; Terazono, K.; Yamamoto, H.; Okamoto, H.
J. Biol. Chem. 265, 7432-7439, 1990
A;Title: Complete nucleotide sequence of human reg gene and its expression i product of the gene.
A;Reference number: A35197; MUID:90237042; PMID:2332435
A;Accession: A35197
                                                                                                                      A;Note: sequence determined from protein isolated after human cDNA sequence was cloned R;Note Caro, A.M.; Adrich, Z.; Fournet, B.; Capon, C.; Bonicel, J.J.; de Caro, J.D.; Rove Biochim. Biophys. Acta 994, 281-284, 1989
A;Title: N-terminal sequence extension in the glycosylated forms of human pancreatic st A;Reference number: S02767; MUID:89150292; PMID:2493268
A;Accession: S02767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: GB:M18963; NID:g190978; PIDN:AAA36558.1; PID:g190979 R;Itoh, T.; Tsuzuki, H.; Katoh, T.; Teraoka, H.; Matsumoto, K.; Yoshida, FEBS Lett. 272, 85-88, 1990 A;Title: Isolation and characterization of human reg protein produced in A;Reference number: S12950; MUID:91032149; PMID:2226837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: UNIPROT:P05451; GB:J05412
R;Terazono, K.; Yamamoto, H.; Takasawa, S.; Shiga, K.; Yamamoto, H.; Takasawa, S.; Shiga, K.; Yamamoto, H.; Takasawa, S.; Shiga, K.; Yamamoto, Takasawa, S.; Shiga, K.; Yamamoto, Ya
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A; Molecule type: protein
A; Residues: 23-47 < DEC>
R; Rouimi, P.; de Caro, J
                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: protein
A; Residues: 23-52; 160-166 < ITO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Accession: S12950
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A; Residues: 1-166 < TER>
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A; Residues: 1-166 <WAT>
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A;Molecule type: mRNA
A;Residues: 1-174 <FRI>
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34.5%; Pred. No. 3.6e-15;
tive 25; Mismatches 57
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de Caro, J.;

Bonicel,

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Caro,

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pancreatic stone protein precursor - human (Species: Homo sapiens (man) (C;Species: Homo sapiens (man) (C;Date: 03.Jun-1993 #sequence_revision 03.Jun-1993 #text_cl (C;Accession: A45751 R;Giorgi, D.; Bernard, J.P.; Rouquier, S.; Iovanna, J.; Saz J. Clin. Invest. 84, 100-106, 1989 A;Title: Secretory pancreatic stone protein messenger RNA. A;Reference number: A45751; MUID:89292148; PMID:2525567
                                                                                                                                                                                                                   RESULT 12
A45751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Map position: 2p12-2p12
A;Introns: 2p11, 61/3; 107/3; 145/1
A;Introns: 2p12, 61/3; 107/3; 145/1
C;Superfamily: tetranectin; C-type lectin homology
C;Keywords: glycoprotein; lectin; pancreas; pyroglutamic acid
F;1-22/Domain: signal sequence #status predicted <SIG>F;1-22/Domain: signal sequence #status predicted <SIG>F;1-22/Domain: signal sequence #status predicted <SIG>F;1-22/Domain: capacreatic stone protein #status experimental <MAT2>F;36-162/Domain: C-type lectin homology <LCH>F;36-162/Domain: C-type lectin homology <LCH>F;36-162/Domain: C-type lectin homology <LCH>F;23/Modified site: pyrrolidone carboxylic acid (Gln) (in mature form) #status
F;27/Binding site: carbohydrate (Thr) (covalent) #status experimental
F;33-34/Cleavage site: Arg-Ile (trypsin) #status experimental
F;36-47,64-162,137-154/Disulfide bonds: #status experimental
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A; Molecule type: protein
A; Residues: 34-73, X', 75-87, 'R', 89-98 < MON>
C; Comment: This protein is found in pancreatic calculi of mammals.
C; Comment: Intact regenerating islet lectin 1-alpha, lithostathine,
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A; Residues: 33-48 < RO2;
R; Montalto, G.; Bonicel, J.; Multigner, L.; Rovery, M.; Sarles,
Biochem. J. 238, 227-232, 1986
A; Title: Partial amino acid sequence of human pancreatic stone F
A; Reference number: A25246; MUID:87099950; PMID:3541906
A; Accession: A25246
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A;Title: Cleavage of the Arg-Ile bond in the native polypeptide chain A;Reference number: S01471; MUID:87219142; PMID:3108036
A;Accession: S01471
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A; Residues: 63-72;125-139;150-157;160-166 < ROU>
A; Residues: 63-72;125-139;150-157;160-166 < ROU>
A; Note: disulfide bonds
R; de Caro, A.M.; Bonicel, J.J.; Rouimi, P.; de Caro,
R; de Caro, B. Biochem. 168, 201-207, 1987
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                                                                                                                                                                                                                                                                                                                                                                                 W--SGKSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKYR 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SYGNGAHLASILSLKEASTIAEYI--SGYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MLISCLMFLSQSQGQEAQTELPQ---ARISCPEGTNAYRSYCYYFNEDRETWVDADLYCQ
                                                                                                                                                                                                                                                                                                                                    WGIGAPSSVNPGYCVSLTSSTGFQKWKDVPCEDKFSFVCKFK
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S02419; MUID:88152214; PMID:3345835
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31.5%; Pred. No. 6.7e-15;
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                                                                                    J.; Sarles,
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                                                                                    H.; Dagorn,
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                                                                                    J.C.
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A;Title: Isolation and charac
A;Reference number: S32489; M
A;Accession: S32489
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-172 <TIF>
pancreatitis-associated protein precursor - human pancreatitis-associated protein precursor - human pancreatic stone protein homol C; Species: Homo sapiens (man) C; Date: 20-Feb-1995 #sequence revision 20-Feb-1995 #text_change C; Accession: A49616; A44931; S29821; S48197; I55880 R; Dusetti, N.J.; Frigerio, J.M.; Fox, M.F.; Swallow, D.M.; Dagori Genomics 19, 108-114, 1994 A; Title: Molecular cloning, genomic organization, and chromosoma A; Title: Molecular cloning, genomic organization, and chromosoma A; Reference number: A49616; MUID:94245143; PMID:8188210 A; Accession: A49616
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C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #
C;Accession: S32489; S28530
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A;Status: preliminary; not compared with conceptual translation
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A;Status: preliminary; not compared with conceptual translation
A;Molecule type: mRNA
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Eur. J. Biochem. 213, 901-907, 1993
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33.3%;
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                                                                                    genomic organization, and chromosomal MUID:94245143; PMID:8188210
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Pred. No. 9.4e-15;
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agkisacutacin alpha chain precursor - sharp-nosed v N;Alternate names: fibrinogenlytic venom protein C;Species: Agkistrodon acutus (bharp-nosed viper) C;Date: 04-Mar-2000 #sequence_revision 04-Mar-2000
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A;Introns: 26/1; 65/3; 111/3; 154/1
C;Superfamily: tetranectin; C-type
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A;Title: Human pancreatitis-associated protein. Messenger RNA A;Reference number: 155580; MUID:93107309; PMID:1469087
A;Accession: I55580
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: GB:D13510; NID:g285970; PIDN:BAA02728.1; PID:g285971 R;Lasserre, C.; Simon, M.T.; Ishikawa, H.; Diriong, S.; Nguyen, V.C.; Christa, Eur. J. Biochem. 224, 29-38, 1994 A;Title: Structural organization and chromosomal localization of a human gene A;Reference number: S48197; MUID:94357229; PMID:8076648 A;Accession: S48197
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A;Title: Cloning and tissue-specific expression of cDNAs
A;Reference number: S29821; MUID:93176807; PMID:7679928
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A;Cross-references: UNIPROT
R;Lasserre, C.; Christa, L.
Cancer Res. 52, 5089-5095,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Cross-references: GDB:136839; OMIM:167805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Gene: GDB: PAP; HIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Status: preliminary; translated from GB/EMBI/DDBJA;Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: mRNA
A; Residues: 1-175 < ITO>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Reference number: S29821;
A; Accession: S29821
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Cross-references: GB:X68641; NID:g312806; PIDN:CAA48605.1; A;Experimental source: hepatocellular carcinoma A;Note: sequence extracted from NCBI backbone (NCBIN:113007, R;Itoh, T.; Teraoka, H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Cross-references: GB:S51768; NID:g262368; PIDN:AAB24642.1; PID:g262369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Residues: 1-175 <LA2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Superfamily: tetranectin, C-type lectin homology; Superfamily: tetranectin, C-type lectin homology; Reywords: acute phase; extracellular protein; pancreas; 1-26/Domain: signal sequence #status predicted <5!G-;27-175/Product: pancreatitis-associated protein #status predicted;40-171/Domain: C-type lectin homology <LCH-;40-51,68-171,146-163/Disulfide bonds: #status predicted
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
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                                                                                                                                                                                                                                                                                                                                                                                                                            MLLSCLMLLSQVQGEEPQRELPSARIRCPKGSKAYGSHCYALFLSPKSWTDADLACQKRP
                                                                                                                                                                                                                                                                                                                         SG-NLVSVLSGAEGSFVSSLVKSIGNSYSYVWIGLHDPTQGTEPNGEGWEWSSSDVMNYF
                                                                                                                                                                                                                                                                  SW--SGKSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKY 156
                                                                                                                                                                                                                 AWERNPSTISSPGHCASLSRSTAFLRWKDYNCNVRLPYVCKF
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5, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25.2%; Score 221; DI 32.7%; Pred. No. 2e-1 tive 29; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2e-14;
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#text_change 09-Jul-2004
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A;Accession: PC7037
A;Molecule type: protein
A;Molecule type: protein
A;Residues: 24-53;84-86;87-94;125-136;137-152 <CH2>
A;Residues: 24-53;84-86;87-94;125-136;137-152 <CH2>
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C;Keywords: disulfide bond; heterodimer; venne
F;11-23/Domain: signal sequence #status predicted <SI
F;24-152/Product: agkisacutacin alpha chain #status
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Accession: JC7134; PC7037
R;Cheng, X.; Qian, Y.; Liu, Q.; Li, B.X.Y.; Zhang, M.; Liu, J. Biochem. Biophys. Res. Commun. 265, 530-535, 1999
A;Title: Purification, characterization, and cDNA cloning of a new A;Reference number: JC7134; MUID:20025379; PMID:10558903
A;Accession: JC7134
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A; Residues: 1-152 < CHE>
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A;Experimental source: venom gland
                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                    Local Similarity
122
                                          126 NKHCAEMSSNNNFLTWSSNECNKROHFLCK 155
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                                                                                                                                                                                                                     10 LLLSCLAKTGVLGDIIMRPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQSYGNGAHLAS
SKKCLGVHIETGFHKWENFYCEQQDPFVCE 151
                                                                                                                                ILSLKEASTIAEYISGYQRSQPI--WIGLHDPQKRQQW--QWIDGAMYLYRSWSGKSMGG 125
                                                                                                                                                                               LLVVFLSLSGTAAD-----CSSGWSSYEGHCYKVFKQSKTWADAESFCTKQVNGGHLVS
                                                                                    IESSGEADFVAHLIAQKIKSAKIHVWIGLRAQNKEKQCSIEWSDGSSISYENWIEEE---
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Pred. No. 4.5e-13;
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Search completed: February 11, 2005, 23:43:48
Job time : 40 secs

Run on:

Searched:

Sequence 1077, Ap Sequence 1078, Ap Sequence 1079, Ap Sequence 138, App Sequence 134, App Sequence 134, App Sequence 136, App Sequence 1200, App Sequence 1200, App Sequence 105, App Sequence 23, Appli Sequence 6394, Appli Sequence 753, App Sequence 113, App Sequence 113, App Sequence 113, Appli Sequence 110, App Sequence 113, Appli Sequence 114, Appli Sequence 27, Appli Sequence 28, Appli Sequence 28, Appli Sequence 38, Appli Sequence 109, App

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Title:
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Copyright (c) 1993 - 2005 Compugen Ltd
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US-09-922-217-1070
US-09-922-217-1077
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Sequence 1070, Ap
Sequence 1077, Ap
Sequence 1078, Ap
Sequence 1080, Ap
Sequence 1080, Ap
Sequence 1077, Ap
Sequence 1077, Ap
Sequence 1078, Ap
Sequence 1079, Ap
Sequence 2078, Ap
Sequence 27, Appli
Sequence 4471, Ap
Sequence 4471, Ap
                                                                                                                                                                                                                                                                                         Description
                                                                                                                                                         APPLICANT: Meagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
APPLICANT: Wang, Tongtong
APPLICANT: Wang, Tongtone Lynn
APPLICANT: King, Gordon E.
APPLICANT: King, Gordon E.
APPLICANT: Clapper, Jonathan D.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
INUMBER OF SEG ID NOS: 1124
SOPTWARE: FastSEQ for Windows Version 4.0
SEG ID NO 1070
LENGTH: 158
TYPE: PRT
ORGANTS"
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US-09-922-217-1070
                                                                                                                                               ; ORGANISM: Homo sapiens
US-09-922-217-1070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 1070, Application US/09922217 Patent No. US/0020076414A1 GENERAL INFORMATION:
                                                           Query Match
Best Local Similarity
Matches 158; Conserva
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1 MASRSMRLLLLLSCLAKTGVLGDIIMRPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQS
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US-10-025-280-1078
US-10-025-380-1079
US-10-025-380-1080
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Database

Result No.

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RESULT 3
US-09-922-217-1078
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US-09-922-217-1077
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                                                                                                                                                                                                                                                                                                         Sequence 1078, Application US/09922217
Patent No. US20020076414A1
GENERAL INFORMATION:
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Best Local Similarity 100.
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APPLICANT: Wang, Aijun
APPLICANT: Clapper, Jonathan D.

TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1077
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APPLICANT:
APPLICANT:
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ORGANISM: Homo sapiens
-09-922-217-1077
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                      Stolk, John A.
Wang, Tongtong
Jiang, Yuqiu
Smith, Carole Lynn
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Meagher, Fac
Meagher, Meagh
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                                                                                                                                           Secrist, Heather
Benson, Darin R.
Meagher, Madeleine
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Smith, Carole Lynn
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Stolk, John A.
Wang, Tongtong
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Benson, Darin R.
King,
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                                                                                                                                                                                                                                                                                  Jiangchun
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Gordon E.
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APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Wang, Tongtong
APPLICANT: Wang, Tongtong
APPLICANT: Ming, Yuqiu
APPLICANT: Ming, Gordon E.
APPLICANT: Ming, Gordon B.
APPLICANT: Wang, Aijun
APPLICANT: Wang, Aijun
APPLICANT: Wang, Aijun
APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, Jonathan D.
FITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USB
FILE REFERENCE: 210121.471C13
CURRENT APPLICATION MUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 11124
COPPEMBER OF SEQ ID NOS: 11124
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                                                                                                                                                                                                                                                       ; TYPE: PRT; ORGANISM: Homo sapiens US-09-922-217-1079
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APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, Jonathan D.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILLING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOPTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1078
LENGTH: 158
LENGTH: 158
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SEQ ID NO 1079
LENGTH: 158
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APPLICANT: Secon
APPLICANT: Benn
APPLICANT: Meac
APPLICANT: Sto
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Best Local Similarity
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YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRSWSG 120
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                                                                                                                                                        100.0%; Score 878; DB 9; Ilarity 100.0%; Pred. No. 3.8e-82; Conservative 0; Mismatches 0;
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YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRSWSG

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APPLICANT: Meagher, Madeleine Joy
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
APPLICANT: Wang, Tongtong
APPLICANT: Smith, Carole Lynn
APPLICANT: Smith, Carole Lynn
APPLICANT: King, Gordon E.
APPLICANT: King, Gordon E.
APPLICANT: Clapper, Jonathan D.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FastSEQ for Windows Version 4.0
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US-09-922-217-1080
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                                                     APPLICANT: Wang, Aijun
APPLICANT: Clapper, Jonathan D.
APPLICANT: Scolk, John A.
APPLICANT: Scolk, John A.
APPLICANT: Meagher, Madelaine J.
APPLICANT: Meagher, Madelaine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121,471C12
FULE REFERENCE: 210121,471C12
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
SOFTWARB: FABLSEQ for Windows Version 3.0
SEQ ID NO 1070
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Best Local Similarity
Matches 158; Conserv
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Patent No. US20020076414A1
GENERAL INFORMATION:
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ORGANISM: Homo sapiens
                                          LENGTH: 158
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Benson, Darin R.
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Sequence 1077, Application US/09833263

Patent No. US20020110547A1

GENERAL INFORMATION:

APPLICANT: Wang, Aljun

APPLICANT: Clapper, Jonathan D.

APPLICANT: Clapper, Jonathan D.

APPLICANT: Clapper, Jonathan D.

TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHER REFERENCE: 210121.471C12

CURRENT APPLICATION NUMBER: US/09/833,263

CURRENT FILING DATE: 2001-04-10

NUMBER OF SEQ ID NOS: 1093

SOFTWARE: FastSEQ for Windows Version 3.0

TYPE: PRI

COLUMNIAN UTTO CRICE
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Sequence 1078, Application US/09833263

Patent No. US20020110547A1

Patent No. US20020110547A1

PAPPLICANT: Wang, Aijun
APPLICANT: Clapper, Jonathan D.
APPLICANT: Stolk, John A.
APPLICANT: Meagher, Madeleine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C12
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
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US-09-833-263-1079
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; ORGANISM: Homo
US-09-833-263-1078
                                                                                                                                                              RESULT 10
                  Sequence 1080, Application US/09833263
Patent No. US20020110547A1
GENERRAL INFORMATION:
APPLICANT: Wang, Aijun
APPLICANT: Clapper, Jonathan D.
APPLICANT: Stolk, John A.
APPLICANT: Meagher, Madeleine J.
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SEQ ID NO 1078
LENGTH: 158
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Best Local
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APPLICANT: Clapper, Jonathan D.
APPLICANT: Stolk, John A.
APPLICANT: Meadher, Madeleine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
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CURRENT FILING DATE: 2001-04-10
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER AND PELLE REFERENCE: 210121.471C12
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ilarity 100.0%;
Conservative 0
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Pred. No. 3.8e-82;
); Mismatches 0;
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APPLICANT: Soppet et al.
TITLE OF INVENTION: Colon Specific Gene and Protein
FILE REFERENCE; PF178D2
CURRENT APPLICATION NUMBER: US/09/525,041
CURRENT FILING DATE: 2001-06-04
PRIOR APPLICATION NUMBER: US 09/162,508
PRIOR FILING DATE: 1998-09-29
PRIOR PPLICATION NUMBER: US 08/468,413
PRIOR APPLICATION NUMBER: US 08/468,413
PRIOR APPLICATION NUMBER: US 08/468,413
PRIOR FILING DATE: 1995-06-06
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn version 3.0
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US-09-525-041-2
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-833-263-1080
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CURRENT FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 1080
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LENGTH: 158
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Best Local Similarity 100.0%;
Matches 158; Conservative 0
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                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                           Best
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                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
                                                                                                                                                                                                                                                       Local
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                    121 KSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKYRP 158
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                                                                            61 YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRSWSG 120
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KSMGGNKHCAEMSSNNNFLTWSSNECNKROHFLCKYRP 158
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Pred. No. 3.8e-82;
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Pred. No. 3.8e-82;
); Mismatches 0;
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RESULT 12
US-09-069-034-4471
; Sequence 4471, Application US/09969034
; Publication No. US20040110668A1

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APPLICANT: Lewis, Marcia E.

TITLE OF INVENTION: Nucleic Acid Sequences Differentially ITILE OF INVENTION: Expressed in Cancer Tissue FILE REFERENCE: 1657/1032

CURRENT APPLICATION NUMBER: US/09/969,034

CURRENT FILING DATE: 2001-10-02

PRIOR APPLICATION NUMBER: 60/237,271

PRIOR FILING DATE: 2000-02-10

NUMBER OF SEQ ID NOS: 4494

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 4471

LEWITH: 158
                                                  APPLICANT: Fanger, GATY R.

APPLICANT: Vedvick Thomas S.

APPLICANT: Vedvick Thomas S.

APPLICANT: Carter, Darrick

IITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS

IITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE

FILE REFERENCE: 210121.471C14

CURRENT APPLICATION NUMBER: US/10/025,380

CURRENT FILING DATE: 2001-12-19

KOUTMARE: FRASESQ for Windows Version 4.0

SEQ ID NO 1070

LENGTH: 158
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US-10-025-380-1070
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Best Local Similarity
TYPE: PRT
ORGANISM: Homo sapiens
-10-025-380-1070
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APPLICANT:
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Carroll, Eddie III
Catino, Theodore J.
Dwlvedi, Poornima
Molino, Gary A.
Thiagalingam, Arunth
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                                                                                                                                                                                                                                                                                    Clapper, Jonathan D. Skeiky, Yasir A. W.
                                                                                                                                                                                                                                                                                                                                                                  Jiang, Yuqiu
Smith, Carole L.
                                                                                                                                                                                                                                                                                                                                                                                                                            Meagher, Madeleine Joy
Stolk, John A.
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Benson, Darin R.
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Wang, Aijun
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APPLICANT: Clapper, Jonathan D.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Vedvick Thomas S.
APPLICANT: Corter, Darrick
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1077
LENGTH: 158
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-025-380-1077
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APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael
APPLICANT: Secrist, Heath
APPLICANT: Benson, Darin
APPLICANT: Meagher, Madel
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Best Local Similarity 100.
Matches 158; Conservative
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APPLICANT:
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Benson, Darin R.
Meagher, Madeleine of
Stolk, John A.
Wang, Tongtong
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                                                                                                         YGNGAHLASILSLKEASTIAEYISGYORSOPIWIGLHDPOKROOWQWIDGAMYLYRSWSG 120
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Smith, Carole L.
King, Gordon B.
Wang, Aijun
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 KSMGGNKHCAEMSSNNNFLTWSSNECNKROHFLCKYRP
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Pred. No. 3.8e-82;
0; Mismatches 0;
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RESULT 15 US-10-025-380-1078 ; Sequence 1078, Application US/10025380 ; Publication No. US20020182191A1

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APPLICANT: Clapper, Jonathan D.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Panger, Gary R.
APPLICANT: Vedvick Thomas S.
APPLICANT: Vedvick Thomas S.
APPLICANT: Vedvick Thomas S.
APPLICANT: Vedvick Thomas S.
APPLICANT: Corter, Darrick
FILE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
FILE OF INVENTION: OP COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1078
LENGTH: 158
TYPE: PRT
ORGANISM: Homo sapiens
Search completed: February 11, 2005, 23:33:33 Job time : 50 secs
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APPLICANT: Ku, Jiangchun
APPLICANT: Lodes, Michael
APPLICANT: Lodes, Michael
APPLICANT: Secrist, Heath
APPLICANT: Secrist, Margher, Madel
APPLICANT: Henson, Darin
APPLICANT: Henson, Darin
APPLICANT: Wang, Tongtong
APPLICANT: Wang, Tongtong
APPLICANT: Smith, Carole
APPLICANT: King, Gordon B
APPLICANT: Wang, Aijun
APPLICANT: Wang, Aijun
APPLICANT: Clapper, Jonat
APPLICANT: Skeiky, Yasir
APPLICANT: Seiky, Yasir
APPLICANT: Carter, Darric
APPLICANT: Carter, Darric
                                                                                                                                                                                                                                                                                                                                                                        Query Match 100.0%; Score 878; DB 13; Best Local Similarity 100.0%; Pred. No. 3.8e-82; Matches 158; Conservative 0; Mismatches 0;
                                                                                   121 KSMGGNKHCAEMSSNNNFLTWSSNECNKROHFLCKYRP 158
                                                                                                                                                                               61 YGNGAHLASILSLKEASTIAEYISGYORSOPIWIGLHDPOKROOWOWIDGAMYLYRSWSG 120
                                                                                                                                                                                                          61 YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAWYLYRSWSG 120
                                                                                                                                                                                                                                                                                                          1 MASRSMRLLLLLLSCLAKTGVLGDIIMRPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQS
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Secrist, Heather
Benson, Darin R.
Meagher, Madeleine Joy
Stolk, John A.
Wang, Tongtong
Jiang, Yuqiu
Smith, Carole L.
King, Gordon E.
Wang, Aijun
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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                            Score
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Gapop 10.0 , Gapext 0.
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878
  513545 seqs, 74649064 residues
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1: /cgn2_6/ptodata/1/1aa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/1aa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/1aa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/1aa/6B_COMB.pep:*

5: /cgn2_6/ptodata/1/1aa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/1aa/backfiles1.pep:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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US-09-621-976-4812
US-08-401-530A-7
US-08-729-103-3
US-08-729-103-3
US-08-729-103-4
US-09-949-016-10686
US-08-401-530A-5
US-08-401-530A-6
US-08-401-530A-6
US-08-401-530A-6
US-08-401-530A-6
US-08-401-530A-6
US-08-949-016-6286
US-09-949-016-6286
US-09-949-016-6286
US-08-464-637-2

US-08-401-530A-4

US-08-709-662-4

US-08-822-261-3

US-09-226-852-3

US-09-949-016-10537

US-07-778-156-7

US-08-822-261-4

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US-08-468-413-2
US-09-162-508-2
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           Sequence 7, Appli
Sequence 3, Appli
Sequence 4, Appli
Sequence 10686, A
Sequence 5, Appli
Sequence 6, Appli
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Sequence 2, Appli
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Sequence 4, Appli
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190	190	193	194.5	194.5	196	196	198.5	198.5	198.5	198.5	204	204	204.5	204.5	208	208	218	
21.6	21.6	22.0		22.2			22.6	22.6	•	22.6		•	23.3	•	23.7	ü	24.8	
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US-08-709-662-2	US-08-401-530A-2	PCT-US95-03747-2	PCT-US92-10344-1	US-07-893-929A-1	PCT-US92-10344-3	US-07-893-929A-3	PCT-US92-10344-2	US-07-893-929A-2	PCT-US92-10344-5	US-07-893-929A-5	US-09-226-852-1	US-08-822-261-1	PCT-US92-10344-7	US-07-893-929A-7	US-08-709-662-3	US-08-401-530A-3	US-09-226-852-4	
•	'n	-	Sequence 1,	۲,	e 3,	Sequence 3,	Sequence 2,	'n	5	<u></u> ა	Sequence 1,	Sequence 1,	Sequence 7,	Sequence 7,	Sequence 3,	Sequence 3,	-	
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ALIGNMENTS

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APPLLICATE
FILING DATE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
NAME: Billings, Lucy J.
NAME: Billings, Lucy J.
NAME: Billings, Lucy J.
NAME: 36,749
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PP-01
TELECHONE: 415-855-0555
TELECHONE: 415-855-0555
TELECHA: 415-845-4166
; INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 158 amino acid
US-08-729-103-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     us-08-729<u>-</u>103<sup>-</sup>1
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Patent No. 38378412
                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOPTWARE: FASTSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/729,103
FILING DATE: Filed Herewith
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                           STRANDEDNESS:
TOPOLOGY: lin
MOLECULE TYPE:
                  IMMEDIATE SOURCE:
LIBRARY: COLNF
CLONE: 1310334
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: BANDMAN, Olga
APPLICANT: Goli, Surya K.
TITLE OF INVENTION: NOVEL HUMAN REG PROTEIN
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U
ZIP: 94304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 3174 Po-
CITY: Palo Alto
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                      1310334
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3174 Porter Drive
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Query Match Best Local Similarity

100.0%; Score 878; DB 2; ilarity 100.0%; Pred. No. 5.9e-88; Conservative 0; Mismatchan

Length 158; Indels

0;

Gaps

0

Matches 158;

MASRSMRLLLLLSCLAKTGVLGDIIMRPSCAPGWFYHKSNCYGYFRKLRNWSDABLECQS

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US-08-468-413-2
                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 158; Conserv
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APPLICANT: LI, ET
                                                                                                                                                                                                                                                                                                                                             TELEPAX: 201-994-1744
INFORMATIÓN FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 158 AMINO ACIDS
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/460
FILING DATE: 06 JUN 95
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION MOMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE/
ATTORNEY AGENTON:
ATTORNEY AGENTON:
NAME: FEREARO, GREGORY D.
REGISTRAȚION NUMBER: 36,134
REGISTRAȚION NUMBER: 3258
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: LI, ET
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN, ADDRESSEE: CECCHI, STEWART & OLSTEIN STREET: 6 BECKER FARM ROAD CITY: ROSELAND STATE: NEW JERSEY
                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
TOPOLOGY: LI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PS,
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
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                   KSMGGNKHCAEMSSNNNFLTWSSNECNKROHFLCKYRP 158
                                                                                   YGNGAHLASILSLKEASTIAEYISGYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRSWSG 120
                                                                                                                                      MASRSMRILLLLSCLAKTGVLGDI IMRPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQS
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KSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKYRP 158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
                                                                                                                                                                                                     100.0%; Score 878; DB 2; Length 158; ilarity 100.0%; Pred. No. 5.9e-88; Conservative 0; Mismatches 0: Tndol-
                                                                                                                                                                                                                                                                                                                   LINEAR
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US-09-162-508-2
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 158 AMINO ACIDS
TYPE: AMINO ACID
                                                                                                                          Sequence 2, Application PC/TUS9507169 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches 158;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/468,41
FILING DATE: 06 JUN 95
ATTORNEY/AGENT INFORMATION:
NAME: FERRARO, GREGORY D.
REGISTRATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 3258
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:

GENERAL INFORMATION:

LI, ET AL.
APPLICANT: LI, ET AL.
TITLE OF INVENTION: Human Colon Specific Gene
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
ADDRESSEE: CEARELLA, BYRNE, BAIN, GILFILLAN,
ADDRESSEE: CECHI, STEWART & OLSTEIN
STREET: 6 BECKER FARM ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/162,508
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TITLE OF INVENTION: HU
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: PROTEIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: ROSELAND
STATE: NEW JERSEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                               121 KSMGGNKHCAEMSSNNNPJÍWSSNECNKRQHFLCKYRP 158
                                                                                                                                                                                                                                                                                                       61
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                                                                                                                                                                                                                                                                                                                                                                                                      1 MASRSMRLILLLSCLAKTGVLGDIIMRPSCAPGWFYHKKNCYGYFRKLRNWSDAELECQS
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ilarity 100.0%;
Conservative (
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Pred. No. 5.9e-88;
); Mismatches 0;
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                                                                                          Gene
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
COMPUTER: IBM PS/2

CITY: ROSELAND STATE: NEW JERSEY COUNTRY: USA

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RESULT 5
US-09-621-976-4812
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                                                                 NUMBER OF SEQ ID NOS: 1935;
SOPTWARE: Patent.pm;
SEQ ID NO 4812;
LENGTH: 122;
TYPE: PRT
ORGANIZM: Homo maplens
US-09-621-976-4812
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                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local Similarity
Query Match
Best Local Similarity
Matches 122; Conserv
                                                                                                                                                                                                                                                                                                                             Sequence 4812, Application US/09621976 Patent No. 6639063
                                                                                                                                                                                  APPLICANT: Giordano, J.Y.

TITLE OF INVENTION: ESTS and Encoded Human Proteins.

FILE REPERIOR: GENSET.054PR2

CURRENT APPLICATION NUMBER: US/09/621,976

CURRENT FILING DATE: 2000-07-21
                                                                                                                                                                                                                                                                                       APPLICANT: Dumas Milne Edwards, J.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 201-994-1700
TELEPAX: 201-994-1744
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                       APPLICANT: Jobert, S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORIEY AGENT INFORMATION:
NAME: FERRARO, GREGORY D.
REGISTRATION NUMBER: 36,134
REFERENCE/DOCKET NUMBER: 325800-389
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US:
FILING DATE: 06 JUN 95
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                KSMGGNKHCAEMSSNNNFLTWSSNECNKROHFLCKYRP 158
 Conservative
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06 JUN 95
              79.2%; Score 695; DB 4; 100.0%; Pred. No. 3.9e-68;
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 Mismatches
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                                 Length 122;
Indels
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US-08-401-530A-7
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                                                                                                                                                                                          Query Match 29.0%; Score 254.5; DB 2; Best Local Similarity 30.6%; Pred. No. 6.6e-20; Matches 52; Conservative 38; Mismatches 51;
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 165 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/401,530A
FILING DATE: 22-FEB-1995
CLASSIFICATION: 800
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein ORIGINAL SOURCE:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 009
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Rosenberg, Lawrence
APPLICANT: Duguid, William P.
TITLE OF INVENTION: INCAP PROTEIN INVOLVED IN PANCREATIC
TITLE OF INVENTION: ISLET NEOGENESIS
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                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
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STREET: 1001 G Street, N.W.
                                                                                                                                                                                                                                                                                       ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
110 GAMYLYRSW-SGKSMGGNK-HCAEMSSNNNFLTWSSNECNKRQHFLCKYR 157
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5. 5834590
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                                                                             SDAELECOSYGNGAHLASILSLKEASTIAEYI--SGYQRSQPIWIGLHDPQKRQQWQWID 109
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                                                       AEADLFCONWNSG-YLVSVLSQAEGNFLASLIKESGTTAAN-VWIGLHDPKNNRRWHWSS 114
                                                                                                                         KYFILLSCL-----MVLSPSQGQEAEEDLPSARITCPEGSNAYSSYCYYFMEDHLSW 56
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                                                                                                                                                                                                                                                                                       Rattus rattus
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                                                                                                                                                                                                                             DB 2;
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                                                                                                                                                                                                                                 Length 165;
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; Sequence 7, Application US/08709662
Patent No. 5840531
; GENERAL INFORMATION:
APPLICANT: Vinik, Aaron I.
APPLICANT: Pittenger, Gary L.
                                                                                         RESULT 8
US-08-709-662-7
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                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
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NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Bandman, Olga
APPLICANT: Goli, Surya K.
TITLE OP INVENTION: NOVEL HUMAN REG PROTEIN
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IMMEDIATE SOURCE:
LIBRARY: GenBa
CLONE: 393209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/7
FILING DATE: Filed Herewith
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc
STREET: 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM CON OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH:
                                                                                                                                                                                                     110 GAMYLYRSW-SGKSMGGNK-HCAEMSSNNNFLTWSSNECNKRQHFLCKYR 157
                                                                                                                                                                                                                                                57 AEADLFCQNMNSG-YLVSVLSQAEGNFLASLIKESGTTAAN-VWIGLHDPKNNRRWHWSS 114
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                                                                                                                                                                                                                                                                                 52 SDAELECQSYGNGAHLASILSLKEASTIAEYI--SGYQRSQPIWIGLHDPQKRQQWQWID 109
                                                                                                                                                                                                                                                                                                                                                                                                   52;
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                                                                                                                                                                                                                                                                                                                                                          7 RLLLLLSCLAKTGVLGDIIMRPS------CAPGWFYHKSNCYGYFRKLRNW 51
                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                                      KYFILLSCL-----MVLSPSQGQEAEEDLPSARITCPEGSNAYSSYCYYFMEDHLSW 56
                                                                                                                                                                      GSLFLYKSWDTGYPNNSNRGYCVSVTSNSGYKKWRDNSCDAQLSFVCKFK 164
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                                                                                                                             29.0%; Score 254.5; DB 2
30.6%; Pred. No. 6.6e-20;
tive 38; Mismatches 51
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                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 165;
                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                 29;
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                                                                                                                                                                                                                                            Sequence 4, Appli
Patent No. 583784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Bandman, Olga
APPLICANT: Bandman, Olga
APPLICANT: GOLI, Surya K.
TITLE OF INVENTION: NOVEL HUMAN REG PROTEIN
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
ORIGINAL SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 165 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOPTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA: APPLICATION UNBER: US/08/709,662 FILING DATE: 09-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Rosenberg, Lawrence
APPLICANT: Duguid, William :
TITLE OF INVENTION: INCAP PROTEIN INVOLVED IN PANCREATIC
TITLE OF INVENTION: ISLET NEOGENESIS
 COMPUTER READABLE FORM:
                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                    STREET: 3174 POI
CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 00
                   COUNTRY: UZIP: 94304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                               ADDRESSEE:
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Local Similarity 30.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Rattus rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
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                                                                                                                                                                                                                                                                                                                                                            115
                                                                                                                                                                                                                                                                                                                                                                                               110 GAMYLYRSW-SGKSMGGNK-HCAEMSSNNNFLTWSSNECNKRQHFLCKYR 157
                                                                                                                                                                                                                                                                                                                                                                                                                                  57 AEADLFCQNMNSG-YLVSVLSQAEGNFLASLIKESGTTAAN-VWIGLHDFKNNRRWHWSS 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     52 SDAELECOSYGNGAHLASILSLKEASTIAEYI -- SGYQRSQPIWIGLHDPQKRQQWQWID 109
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                                                                                                                                                                                                                                                                                                                                                          GSLFLYKSWDTGYPNNSNRGYCVSVTSNSGYKKWRDNSCDAQLSFVCKFK 164
                                                                                                                                                                                                                                                                Application US/08729103
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                                                                                         E: Incyte Pharmaceuticals, 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B: Banner & Witcoff, Ltd.
1001 G Street, N.W.
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)N: 435
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Pred. No. 6.6
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DB 2; 51;

Length 165; Indels 29;

Gaps

OPERATING SYSTEM: DOS SOFTWARE: FastSEQ Version 1.5

: Diskette IBM Compatible YSTEM: DOS

COMPUTER: II

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                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR PILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
                                                                                                              SEQ ID NO 10686
LENGTH: 174
TYPE: PRT
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Best Local !
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: VENTER, J.
                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                   ORGANISM: Human
                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 4:
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NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                       NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
IMMEDIATE SOURCE:
LIERARY: GenBank
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 166 amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'Match 27.3%; Score 240; DB 2; Length 166; Local Similarity 34.3%; Pred. No. 2.6e-18; Local Similarity 27; Mismatches 57; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLONE: 474306
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QRSQP-IWIGIHDPQKRQQWQWIDGAMYLYRSW-SGKSMGGNK-HCAEMSSNNNFLTWSS 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RISCPEGTNAYRSYCYYFNEDPETWVDADLYCQNMNSG-NLVSVLTQAEGAFVASLIKES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STDDSNVWIGLHDPKKNRRWHWSSGSLVSYKSWDTGSPSSANAGYCASLTSCSGFKKWKD 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    166 amino acids
  27.3%;
34.3%;
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Score 240; DB 4; Length 174; Pred. No. 2.7e-18;
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US-08-401-530A-5
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APPLICANT: Vinik,
APPLICANT: Pitteng
APPLICANT: Rafaelo
APPLICANT: Rosenbe
APPLICANT: Duguid,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                         Matches
                                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 202-508-9299
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 174 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 005
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/401,530A
FILING DATE: 22-FEB-1995
CLASSIFICATION: 800
                                                                                                                                                                                                                                                           TOPOLOGY: linear MOLECULE TYPE: protein ORIGINAL SOURCE: ORGANISM: Rattus rati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 20001-4597
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Duguid, William P.
TITLE OF INVENTION: INGAP PROTEIN INVOLVED IN PANCREATIC
TITLE OF INVENTION: ISLET NEOGENESIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
CITY: W
                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Banner & Allegretti
STREET: 1001 G Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                160 ESCEKKFSFVCKFK 173
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65 DLACQKRPSG-HLVSVLSGSEASFVSSLIKSSGNSGQNVWIGLHDPTLGQEPNRGGWEWS 123
                                     55 ELECQSYGNGAHLASILSLKEASTIAEYI-SGYQRSQPIWIGLHDPQKRQQ-----WQWI 108
                                                                                                                                                           55;
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                                                                           VALTTMSWMLLSSLMLLSQVQGEDAKEDVPTSRISCPKGSRAYGSYCYALFSVSKSWFDA
                                                                                                               MASRSMRLLLLLSCLAKTGVLGD-----IIMRPSCAPGWFYHKSNCYGYFRKLRNWSDA
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                                                                                                                                                                                                                                                                                                                                                           174 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                        202-508-9299
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Rafaeloff, Ronit
                                                                                                                                                      26.3%; Score 231; DB 2; Length 174; ilarity 32.4%; Pred. No. 2.6e-17; Conservative 28; Mismatches 73; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Rosenberg,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vinik, Aaron I.
                                                                                                                                                                                                                                                           Rattus rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lawrence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     00570.48743
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                                                                                                                                                           Indels 14;
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                                                                               64
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RESULT 12
US-08-709-662-5
; Sequence 5, Application No. 5840531
RESULT 13
US-08-401-530A-6
i Sequence 6, Application US/08401530A
; Patent No. 5834590
; GENERAL INFORMATION:
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US-08-709-662-5
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 202-508-9299
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 174 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/709,662
FILING DATE: 09-SEP-1996
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
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APPLICANT: Rafaeloff, Ronit
APPLICANT: Rosenberg, Lawrence
APPLICANT: Duguid, William P.
TITLE OF INVENTION: INGAP PROTEIN INVOLVED IN PANCREATIC
TITLE OF INVENTION: ISLET NEOGENESIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 1001 G SI
CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: D.C
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STREET: 1001 G Street, N.W.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           109
                                                                                                                                               124 NADVMNÝFNWETNPSSVSGSHĆGTLTRASGFLRWRENNĆISELPYVĆKFK 173
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                                                                                                                                                                                                                                                                                                                                       1 MASRSMRLLLLLSCLAKTGVLGD-----IIMRPSCAPGWFYHKSNCYGYFRKLRNWSDA
                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
)GY: linear
                                                                                                                                                                                 DGAMYLYRSW-SGKSMGGNKHCAEMSSNNNFLTWSSNECNKRQHFLCKYR 157
                                                                                                                                                                                                                                                 BLECQSYGNGAHLASILSIKEASTIAEYI-SGYQRSQPIWIGLHDPQKRQQ-----WQWI 108
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                                                                                                                                                                                                                       DLACOKRPSG-HLVSVLSGSEASFVSSLIKSSGNSGONVWIGLHDPTLGQEPNRGGWEWS
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                                                                                                                                                                                                                                                                                                                                                                           26.3%; Score 231; DB 2; Length 174; 32.4%; Pred. No. 2.6e-17; tive 28; Mismatches 73; Indels
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                                                             Sequence 6, Application US/08709662
Patent No. 5840531
GENERAL INFORMATION:
APPLICANT: Vinik, Aaron I.
APPLICANT: Pittenger, Gary L.
APPLICANT: Rafaeloff, Ronit
APPLICANT: Rafaeloff, Ronit
Duguid, William P.
TITLE OF INVENTION: INGAP PROTE
TITLE OF INVENTION: INGAP PROTE
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US-08-709-662-6
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Version #1.25
SOFTWARE: PAtentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/401,530A
FILING DATE: 22-FEB-1995
CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: Kagan, Sarah A.
REGISTRATION NUMBER: 32,141
REGISTRATION NUMBER: 32,141
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Best Local Similarity
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ORIGINAL SOURCE:
ORGANISM: Ratt
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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ZIP: 20001-4597
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Rosenberg, Lawrence
APPLICANT: Duguid, William P.
TITLE OF INVENTION: INGAP PROTEIN INVOLVED IN PANCREATIC
TITLE OF INVENTION: ISLET NEOGENESIS
NUMBER OF SEQUENCES: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS
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                             NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: D.C
             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               36 RTSCPMGSKAYRSYCYTLVTTLKSWFQADLACQKRPSG-HLVSILSGGEASFVSSLVTGR
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                                                                                                                                                                                                                                                                                                                                       LKWGDHHCDVELPFVCKFK 173
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1001 G Street, N.W.
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Rafaeloff, Ronit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vinik, Aaron I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rattus rattus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                202-508-9100
           Banner & Witcoff, Ltd.
                               ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein
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                                                   INGAP PROTEIN INVOLVED IN PANCREATIC ISLET NEOGENESIS
                                                                                                                             Lawrence
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Pred. No. 4.9e-17
25; Mismatches 5
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Indels Length

9

Gaps

STREET: CITY: W

Washington

1001 G Street, N.W.

D.C.

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; ORGANISM: Human US-09-949-016-10685
                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 26.0%; Score 228.5; DB 2; Length Best Local Similarity 34.5%; Pred. No. 4.9e-17; Matches 48; Conservative 25; Mismatches 57; Indels
                                                              NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 10685
LENGTH: 144
                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 10685, Application US/09949016 Patent No. 6812339
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FILING DATE: 09-SEP-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: KAGGAT, SETAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 0057
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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INFORMATION FOR SEQ ID NO: 6:
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ZIP: 20001-4597
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION NUMBER: US/08/709,662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein ORIGINAL SOURCE: ORGANISM: Rattus ratt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 174 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 174 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 202-508-9299
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              86 YORSOPIWIGLHDPOKROO-----WOWIDGAMYLYRSWSG--KSMGGNKHCAEMSSNNNF 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              36 RTSCPMGSKAYRSYCYTLVTTLKSWFQADLACOKRPSG-HLVSILSGGEASFVSSLVTGR
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Search completed: February 11, 2005, 23:43:03
Job time : 44 secs
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                                                                                129 DVPCEDKFSFVCKFK 143
                                                                                                                   143 SNECNKROHFLCKYR 157
                                                                                                                                                                                 85 GYQRSQPIWIGLHDPQKRQQWQWIDGAMYLYRSW--SGKSMGGNKHCAEMSSNNNFLTWS 142
                                                                                                                                                                                                                                     27 RPSCAPGWFYHKSNCYGYFRKLRNWSDAELECQSYGNGAHLASILSLKEASTIAEYI--S
                                                                                                                                                          70 GTDDFN-VWIGLHDPKKNRRWHWSSGSLVSYKSWGIGAPSSVNPGYCVSLTSSTGFQKWK 128
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11112 bade Blank (nepto)